SEQUENCE LISTING

<110> Aros Applied Biotechnology ApS

<120> Classification of Colon Cancer

<130> P949PC00

<160> 139

<170> PatentIn version 3.1

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| acatttcgag | atgaccttct | caatttgcta | agagaaagcc | gatttgactt | tatctacgat | 1980 |
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<211> 3453

<212> DNA

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| aagcatgtag | actgtgccag | aggccagacc | cacgggctca | tgcacccctg | agccagcagg | 2520 |
| gcatcttgga | aaaggaactc | ttggttcgat | acctggagca | gaggagggga | aagtccaggg | 2580 |
| ctatagggtg | tgatgaagtc | acccctttct | gtcccactac | atctgggact | gactttccga | 2640 |
| gcctccagtc | caaagccggc | ttgatttccg | tgaactctgg | tgctcctgca | tctcatgagt | 2700 |
| gtgccccatg | ggtcccctcc | cctctcagca | tttccttgtc | ccgtctggac | ctggggagtg | 2760 |
| gttaggcagc | aagctttggt | ttatggtttt | cattcattgg | tgaagtaaat | taggcagtgc | 2820 |
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| taaagcctgt | gggtttggtc | cttgaacaag | atgtgggcct | tgcaagatgg | gagagtaaac | 2880 |
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| cttgaagggc | tttattaaag | aaataaaaaa | gaacttttgt | atcttttatc | ctgggagcac | 2940 |
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| taaggatttg | aaccatgaag | tcataggtta | cagacctcag | ttttatgccc | cattggatta | 3120 |
| ctttttttt | tttttttt | tttttttt | tactctttga | aagctttgtt | ttgtggtagt | 3180 |
| ccttttggga | agaatccagt | attatctaca | attattggca | aagtttaaat | gtattttaca | 3240 |
| taacggaaag | tttttagaat | gttgaaaagt | aattgaaaaa | ggtgataggt | aaatttttag | 3300 |
| gcaaagataa | tttatttcaa | taaatctttc | aaaagcctta | ccttgaaatg | ctgttagtaa | 3360 |
| atttctgtga | tttttttt | taatttgttt | tgctgagagc | atagctattt | gtttttattg | 3420 |
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<210> 13

<211> 1351

<212> DNA

 $<\!\!213\!\!>$ NM_013974.1| Homo sapiens dimethylarginine dimethylaminohydrolase 2 (DDAH2), mRNA

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| 60 | cgtagtatat | cctgccccat | ccgtcgcgcc | agccgccaga | aatgccccgg | <400> 13 ccgcttagac |
| 120 | gaggcttgaa | cccagtcccc | agccagagct | ccccgctaaa | acacaaggac | gagctcgcct |
| 180 | gccgagatca | tggggcccca | cctcgggggg | ccaactctgt | cccttctcca | gacggggact |
| 240 | aaactaagaa | agaaacaaga | agacaggtga | tggccgctgg | ggagtggggg | cagcgcgaca |
| 300 | ggaggggctg | ggacgccggg | gatgggatgg | agtctgtgtg | ttggaggggg | atccgagcgg |
| 360 | gggggaaggt | gcctggcgtc | gtcccagaga | gatccgggga | cccatgccct | ggccgctgct |
| 420 | cggggtgctg | aaagggagca | gccaaagctc | tctggatctg | gccttcccgc | gcgggggctg |
| 480 | tgaggagtca | aactgccacc | cagctgctag | actggggcta | tgaggcaacg | ggaggtaaac |
| 540 | ggccctaatc | aaggggacac | gccgtgatcc | tggcgacacg | gaccgctgct | ttgccgctgg |
| 600 | agccctgcaa | gagtccgcaa | gaggtcgatg | tcgtaggcca | ggagccccgc | acgcggccct |
| 660 | tggcactgac | cgacgctgga | gacgagaacg | ggaaatagga | tccgaattgt | gacctggggc |
| 720 | tcaccgagga | aatggaccaa | ggcctctcca | gtttttcgta | ccggccggga | gttctcttca |
| 780 | agtctcgggt | ccactgtgcc | ttcgccgtct | gttccgggac | tggcggacac | gctgagatcg |
| 840 | ggcaggcagc | gcactgttgt | gggggacctc | ctgcggcatg | tgcgcggtct | ccctcccacc |
| 900 | cccatatgcc | tgacagatca | atggcagtgc | tgtccgggca | cccaaaaggc | agcgacgctg |
| 960 | tgggttgcct | ttcttcgtcc | gactgcctct | cgcagctgct | tcccagatga | tccctgaccc |

| ggtgtgcccc | ctttcctcct | gcaccgtgga | ggtggggatc | tgcccaacag | ccaggaggca | 1020 |
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| ggccttgggg | tactgctggc | caggggtagg | atagtatagg | aagtagaagg | ggaaggaggg | 1200 |
| ttagatagag | aatgctgaat | aggcagtagt | tgggagagag | cctcaatatt | gggggagggg | 1260 |
| agagtgtagg | gaaaaggatc | cactgggtga | atcctccctc | tcagaaccaa | taaaatagaa | 1320 |
| ttgacctttt | aaaaaaaaa | aaaaaaaaa | a | | | 1351 |

<210> 14

<211> 4180

<212> DNA

<213> NM_006291.2| Homo sapiens tumor necrosis factor, alpha-induced protein 2 (TNFAIP2), mRNA

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| cagaggcctc | catgtcggag | gcctcctctg | aggacctggt | gccacccctg | gaggctgggg | 180 |
| cagccccata | tagggaggag | gaagaggcgg | cgaagaagaa | gaaggagaag | aagaagaagt | 240 |
| ccaaaggcct | ggccaatgtg | ttctgcgtct | tcaccaaagg | gaagaagaag | aagggtcagc | 300 |
| ccagctcagc | ggagcccgag | gacgcagccg | ggtccaggca | ggggctggat | ggcccgcccc | 360 |
| ccacagtgga | ggagctgaag | gcggcgctgg | agcgcgggca | gctggaggcg | gcgcggccgc | 420 |
| tgctggcgct | ggagcgggag | ctggcggcgg | cggcggcggc | gggcggtgtg | agcgaggagg | 480 |
| agctggtgcg | gcgccagagc | aaggtggagg | cgctgtacga | gctgctgcgc | gaccaggtgc | 540 |
| tgggcgtgct | gcggcggccg | ctggaggcgc | cgcccgagcg | gctgcgccag | gcgctggccg | 600 |
| tggtggcgga | gcaggagcgc | gaggaccgcc | aggcggcggc | ggcggggccg | gggacctcgg | 660 |
| ggctggcggc | cacgcgcccg | cggcgctggc | tgcagctgtg | gcggcgcggc | gtggcggagg | 720 |
| cggccgagga | gcgcatgggc | cagcggccgg | ccgcgggcgc | cgaggtcccc | gagagcgtct | 780 |
| ttctgcactt | gggccgcacc | atgaaggagg | acctggaggc | cgtggtggag | cggctgaagc | 840 |
| cgctgttccc | cgccgagttc | ggcgtcgtgg | cggcctacgc | cgagagctac | caccagcact | 900 |
| tcgcggccca | cctggccgcc | gtggcgcagt | tcgagctgtg | cgagcgcgac | acctacatgc | 960 |
| tgctgctctg | ggtgcagaac | ctctacccca | atgacatcat | caacagcccc | aagctggtgg | 1020 |
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| agctagaggc | acggcgctgg | gctgaggatg | tgcctcccca | gaggctggac | ggccactgcc | 1200 |

| acagcgagct | ggccatcgac | atcatccaga | tcacctccca | ggcccaggcc | aaggccgaga | 1260 |
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| tcctgaggag | ctaccagcgc | gcctttaatg | aatttctgga | gagaggcaag | cagctgacga | 1380 |
| attacagggc | caatgttatt | gccaacatca | acaactgcct | gtccttccgg | atgtccatgg | 1440 |
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| gcctggtcct | caagacggcc | gagcagcagc | agcagctggc | tgggtacatc | ctggccaatg | 1800 |
| ctgacaccat | ccagcacttc | tgcacccagc | acggctcccc | ggcgacctgg | ctgcagcctg | 1860 |
| ctctccctac | gctggccgag | atcattcgcc | tgcaggaccc | cagtgccatc | aagattgagg | 1920 |
| tggccactta | tgccacctgc | taccctgact | tcagcaaagg | ccacctgagc | gctatcctgg | 1980 |
| ccatcaaggg | gaacctatcc | aacagtgagg | tcaagcgcat | ccggagcatc | ttggacgtca | 2040 |
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| tgcctccctc | caccactttt | tttttttt | ttttgagaca | gggtcttgct | gtgttgccca | 3300 |
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| tataataata | attattttt | agagtactgc | ttttgtatgt | atgttgaaca | ggatccaggt | 4140 |
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<210> 15

<211> 2524

<212> DNA

<213> NM_000249.2| Homo sapiens mutL homolog 1, colon cancer, nonpolyposis type 2 (E. coli) (MLH1), mRNA

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| gtagctgatg | ttaggacact | acccaatgcc | tcaaccgtgg | acaatattcg | ctccatcttt | 720 |
|------------|------------|------------|------------|------------|------------|------|
| ggaaatgctg | ttagtcgaga | actgatagaa | attggatgtg | aggataaaac | cctagccttc | 780 |
| aaaatgaatg | gttacatatc | caatgcaaac | tactcagtga | agaagtgcat | cttcttactc | 840 |
| ttcatcaacc | atcgtctggt | agaatcaact | tccttgagaa | aagccataga | aacagtgtat | 900 |
| gcagcctatt | tgcccaaaaa | cacacaccca | ttcctgtacc | tcagtttaga | aatcagtccc | 960 |
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| agcatcctgg | agcgggtgca | gcagcacatc | gagagcaagc | tcctgggctc | caattcctcc | 1080 |
| aggatgtact | tcacccagac | tttgctacca | ggacttgctg | gcccctctgg | ggagatggtt | 1140 |
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| gctgccaaaa | atcagagctt | ggagggggat | acaacaaagg | ggacttcaga | aatgtcagag | 1440 |
| aagagaggac | ctacttccag | caaccccaga | aagagacatc | gggaagattc | tgatgtggaa | 1500 |
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| taccagatac | tcatttatga | ttttgccaat | tttggtgttc | tcaggttatc | ggagccagca | 1800 |
| ccgctctttg | accttgccat | gcttgcctta | gatagtccag | agagtggctg | gacagaggaa | 1860 |
| gatggtccca | aagaaggact | tgctgaatac | attgttgagt | ttctgaagaa | gaaggctgag | 1920 |
| atgcttgcag | actatttctc | tttggaaatt | gatgaggaag | ggaacctgat | tggattaccc | 1980 |
| cttctgattg | acaactatgt | gccccctttg | gagggactgc | ctatcttcat | tcttcgacta | 2040 |
| gccactgagg | tgaattggga | cgaagaaaag | gaatgttttg | aaagcctcag | taaagaatgc | 2100 |
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| ttctgatagt | attcctttat | acacagtgga | ttgattataa | ataaatagat | gtgtcttaac | 2520 |
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<210> 16
<211> 1536
<212> DNA
<213> NM_001071.1| Homo sapiens thymidylate synthetase (TYMS), mRNA
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| caaaaacatg | tatgtgcatt | tcaatcccac | gtacttataa | agaaggttgg | tgaatttcac | 1380 |
| aagctatttt | tggaatattt | ttagaatatt | ttaagaattt | cacaagctat | tccctcaaat | 1440 |
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25

<210> 17

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| cattcattaa | aaagaaacca | ggaaacactt | ttgaaataac | agcataagga | acttcactgt | 1500 |
| ctctgctcaa | taaaatacct | gtaactggaa | aaaaaaaaa | aaaaaaaaa | aaaaaa | 1556 |
| | | | | | | |

<210> 21

<211> 1276

<212> DNA

<213> NM_003581.1| Homo sapiens NCK adaptor protein 2 (NCK2), mRNA

| <400> 21 gtgccaaaga | aggactccat | gaaagatgac | agaagaagtt | attgtgatag | ccaagtggga | 60 |
|------------------------|------------|------------|------------|------------|------------|-----|
| ctacaccgcc | cagcaggacc | aggagctgga | catcaagaag | gtgaacgagc | ggctgtggtt | 120 |
| gctggacgac | tccaagacgt | ggtggcgggt | gaggaacgcg | gccaacagga | cgggctatgt | 180 |
| accgtccaac | tacgtggagc | ggaagaacag | cctgaagaag | ggctccctcg | tgaagaacct | 240 |
| gaaggacaca | ctaggcctcg | gcaagacgcg | caggaagacc | agcgcgcggg | atgcgtcccc | 300 |
| cacgcccago | acggacgccg | agtaccccgc | caatggcagc | ggcgccgacc | gcatctacga | 360 |
| cctcaacato | ccggccttcg | tcaagttcgc | ctatgtggcc | gagcgggagg | atgagttgtc | 420 |
| cctggtgaag | gggtcgcgcg | tcaccgtcat | ggagaagtgc | agcgacggtt | ggtggcgggg | 480 |
| cagctacaac | gggcagatcg | gctggttccc | ctccaactac | gtcttggagg | aggtggacga | 540 |

| ggcggctgcg | gagtccccaa | gcttcctgag | cctgcgcaag | ggcgcctcgc | tgagcaatgg | 600 |
|------------|------------|------------|------------|------------|------------|------|
| ccagggctcc | cgcgtgctgc | atgtggtcca | gacgctgtac | cccttcagct | cagtcaccga | 660 |
| ggaggagctc | aacttcgaga | agggggagac | catggaggtg | attgagaagc | cggagaacga | 720 |
| ccccgagtgg | tggaaatgca | aaaatgcccg | gggccaggtg | ggcctcgtcc | ccaaaaacta | 780 |
| cgtggtggtc | ctcagtgacg | ggcctgccct | gcaccctgcg | cacgccccac | agataagcta | 840 |
| caccgggccc | tcgtccagcg | ggcgcttcgc | gggcagagag | tggtactacg | ggaacgtgac | 900 |
| gcggcaccag | gcgcagtgcg | ccctcaacga | gcggggcgtg | gagggcgact | tcctcattag | 960 |
| ggacagcgag | tcctcgccca | gcgacttctc | cgtgtccctt | aaagcgtcag | ggaagaacaa | 1020 |
| acacttcaag | gtgcagctcg | tggacaatgt | ctactgcatt | gggcagcggc | gcttccacac | 1080 |
| catggacgag | ctggtggaac | actacaaaaa | ggcgcccatc | ttcaccagcg | agcacgggga | 1140 |
| gaagctctac | ctcgtcaggg | ccctgcagtg | acggcgcccc | ggccccacac | tcgcctcccg | 1200 |
| ggccccacgg | tggagctgcc | cgcccggcct | tgtggcagag | gctcctcccg | cggggacggc | 1260 |
| cccgacggct | tctctg | | | | | 1276 |
| | | | | | | |

<210> 22

<211> 1577

<212> DNA

<213> NM_006214.2| Homo sapiens phytanoyl-CoA hydroxylase (Refsum disease) (PHYH), mRNA

| | | | | | | 400 00 |
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| 120 | ggccgcccct | gggccacctc | agattgttct | gcccgtctgc | tcgcgccgcc | tggagcagct |
| 180 | agtttccatc | ttcctctgcc | cagggactat | catcccactt | tgtcgtagct | cggccggggc |
| 240 | agaaaatttt | cctggaacag | atgttctaac | ctggataata | ccagtatact | ctcaacaatt |
| 300 | attcaacgct | tgatgccgat | atcttgtacc | gtaatcaaaa | tgggtttcta | atgaagaaaa |
| 360 | ttaacagtaa | accattagga | aggaggtgaa | atctgcagaa | gtttgaaaaa | ttcggaatga |
| 420 | atcacgaagg | tgagaagatg | atgctccaag | aaatccgaat | gaccatttcg | tgagagatgt |
| 480 | gagattctga | cactctcccc | tcagatactg | aaggagctct | ccaggaagat | tccaggattt |
| 540 | ttgataaaca | gcacacaatg | ttatggccat | ggacctaata | gtgcttcact | aatatgtgga |
| 600 | ctgcactatt | gcaccaggac | gtcaccccct | aagacgtccc | ttctggcaag | aacctccaga |
| 660 | cacatcagcc | ggcgatggag | gcgcctggac | ctcatcgttt | gcccagcgat | tccccttcag |
| 720 | aagccccacg | gggctccctg | gcacacacaa | gtgctcccag | ctgtctggtt | ggaacaacgg |
| 780 | gactacgagg | cgggatccag | aaatgttcca | ggagttaaca | gtgggagggg | attaccccaa |
| 840 | ttccatcctt | cactgttttc | agaagggcga | ctggtgatgg | ccgggtgcac | aaaacaaggc |

| tgctcatcca | cggatctggt | cagaataaaa | cccagggatt | ccggaaggca | atttcctgcc | 900 |
|------------|------------|------------|------------|------------|------------|------|
| atttcgccag | tgccgattgc | cactacattg | acgtgaaggg | caccagtcaa | gaaaacatcg | 960 |
| agaaggaagt | tgtaggaata | gcacataaat | tctttggagc | tgaaaatagc | gtgaacttga | 1020 |
| aggatatttg | gatgtttcga | gctcgacttg | tgaaaggaga | aagaaccaat | ctttgaaata | 1080 |
| gccatctgct | ataactcttt | caacagaaaa | ccaaaaccaa | acgaaatgtc | taaggaaaat | 1140 |
| gttttcttaa | tgagatgatg | taaccttttc | tatcacttgt | taaaagcaga | aaacatgtat | 1200 |
| caggtactta | attgcataga | gttagttttg | cagcacaatg | gtgttgcttt | aatggaaaaa | 1260 |
| aaaaacagta | aaagtgaaat | attactgttt | taaggaaaac | taatttaggg | tggcagccaa | 1320 |
| taaaggtggt | tggtgtctaa | tttaagtgtt | aaatcaattt | ctttcattca | gttagctctt | 1380 |
| tacccaagaa | gaagtgaatg | atttggagct | tagggtatgt | tttgtatccc | ctttctgata | 1440 |
| aacccattcc | ctaccaattt | tatgtcataa | gagattttt | tccccaaat | ctagaacaat | 1500 |
| gtataataca | ttcacatcta | gtcaagggca | taggaacggt | gtcatggagt | ccaaataaag | 1560 |
| tggatattcc | tgctcgg | | | | | 1577 |
| | | | | | | |

<210> 23

<211> 3060

<212> DNA

<213> NM_004739.2| Homo sapiens metastais-associated gene family, member 2 (MTA2), mRNA

| | | | | | | 400 22 |
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| 120 | ggcctggcag | gcctggctct | ggtagctgga | cgggggcggg | tgggactccg | caacgactag |
| 180 | gtcccgggag | cagcctcagc | ggacgggggc | gaagccgagc | tgttccggaa | gagccgagct |
| 240 | gctgaggcag | aaccgcgacg | ggccgggatg | cgacagcgcg | gctgcggcgg | tgaggcgata |
| 300 | ccaccgttcg | aagcggcagc | gactccctcg | gccccagtga | ggctgcgcgg | cggaggtgcc |
| 360 | tgtgcggccg | cggacccctt | gcgagcctcc | ccctgccccc | tcgagccgag | gggctttgcc |
| 420 | ttacgtctat | gggtgggaga | aacatgtacc | catggcggcc | cgggaacggc | gaggcggcgg |
| 480 | caacaagact | ttgaggagct | gttagacgga | tccttacctg | cttccagcaa | tttgagaact |
| 540 | catttctagt | ggcgcaggga | tgtcttttcc | aaaggttgtc | atgtggaggc | gcaaatggaa |
| 600 | aaa gc agcca | aagaggaatc | agggagtttg | tagtaatgcc | gcctggctga | agcctcaaca |
| 660 | ttctcggcaa | aactttttct | aagcaccggg | ccatcaactg | agcagcagcg | ggggtgtctg |
| 720 | cttgaatgag | gtgtgaccct | gggaaatgca | ccacatacgg | taccagccac | tttgaatcat |
| 780 | actggtgttt | ttttttactc | gaggactgct | cctggaaaag | tgagccagta | acagatatct |
| 840 | ttgcaaatac | ttagagttgg | cagggcgaga | tctcgctgat | agaagacact | gaccccgtgc |

| caagctgaga | tcccagatcg | cctagtagag | ggagaatctg | ataatcggaa | ccagcagaag | 900 |
|------------|------------|------------|------------|---------------------|------------|------|
| atggagatga | aggtctggga | cccagacaac | cctctcacag | accggcagat | cgaccagttt | 960 |
| cttgtggtgg | cccgagctgt | gggaaccttt | gcaagagccc | tagattgtag | cagctccatt | 1020 |
| cggcagccaa | gcttgcacat | gagtgcagct | gctgcctccc | gagatatcac | tctgtttcac | 1080 |
| gccatggata | ccttgcaaag | gaacggctac | gacctggcta | aggccatgtc | gaccctggta | 1140 |
| ccccagggag | gcccggtgct | gtgtcgggat | gagatggagg | aatggtcagc | ctcagaggcc | 1200 |
| atgctatttg | aggaggccct | agagaagtat | gggaaggact | tcaatgatat | tcgccaggat | 1260 |
| tttctaccct | ggaagtcact | tgccagcata | gtccagtttt | attacatgtg | gaaaaccaca | 1320 |
| gaccggtata | ttcagcagaa | aaggttgaaa | gctgctgaag | cagacagcaa | actgaaacag | 1380 |
| gtctacattc | ccacctacac | taagccaaac | cctaaccaga | tcatttctgt | gggttcaaaa | 1440 |
| cctggcatga | atggggctgg | atttcagaag | ggcctgactt | gtgagagttg | ccacaccaca | 1500 |
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| actcggggca | ccacggagcc | acactcaagg | ggtcatttat | ccagacctga | agctcaaagt | 1680 |
| ctctctctt | acacaaccag | cgccaacagg | gccaagctac | tggctaagaa | cagacaaact | 1740 |
| ttcctgcttc | agaccacaaa | gctgacccgt | cttgccagac | gcatgtgcag | ggacctatta | 1800 |
| cagccaagga | gggccgcccg | acggccttat | gctcctatca | atgccaatgc | catcaaagca | 1860 |
| gagtgctcca | ttcgacttcc | taaggccgcc | aagactccat | tgaagattca | ccctctggtg | 1920 |
| cggctgcccc | tggcaactat | cgtcaaagat | ctggtggccc | aggcacccct | gaaaccaaaa | 1980 |
| acacctcggg | gtaccaagac | accgatcaac | agaaaccagc | tgtcccagaa | ccggggactg | 2040 |
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| cagaaactaa | acccagctga | tgcccccaat | cctgtggtgt | ttgtggccac | aaaggatacc | 2220 |
| agggccctac | ggaaggctct | gacccatctg | gaaatgcggc | gagctgctcg | ccgacccaac | 2280 |
| ttgcccctga | aggtgaagcc | aacgctgatt | gcagtgcggc | cccctgtccc | tctacctgca | 2340 |
| ccctcacatc | ctgccagcac | caatgagcct | attgtcctgg | aggactgagc | acctgtgggg | 2400 |
| aagggaggtg | ggctgagagg | tagagggtgg | atgcccaggg | cacccaaacc | tcccttccct | 2460 |
| ttcgtgtcga | agggagtgag | gagtgaatta | aggaagagag | caagtgagtg | tgtgtccctg | 2520 |
| gaggggttgg | gcgccctctg | gtgttaccac | ctcgagactt | gtctcatgcc | tccatgcttg | 2580 |
| ccgatggagg | acagactgca | ggaacttggc | ccatgtggga | acctagcctg | ttttgggggg | 2640 |
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| ttgcctccct | ttgtgaaagg | ggatggggag | gggaagagta | aacagataac | aggtggtggt | 2760 |
| acctggttgg | gggaggggg | cgtgcactgc | catgtctttt | tttttttt | tttttttt | 2820 |
| | | | | | | |

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<210> 24

<211> 2407

<212> DNA

<213> NM_001091.1| Homo sapiens amiloride binding protein 1 (amine oxidase (copper-containing)) (ABP1), mRNA

<400> 24 60 120 tggagcgaga gatgccggcc ctgggctggg ccgtggctgc catcctgatg ctgcagacgg 180 ccatggcgga gccctccccg gggactctgc ccaggaaggc aggggtgttt tcagacctaa 240 qcaaccaaga gctgaaggca gtgcacagct tcctctggtc caagaaggag ctgaggctgc agccctccag taccaccacc atggccaaga acaccgtgtt tctcatcgag atgctgctgc 300 ccaaqaaqta ccatgtgctg aggtttctgg ataaaggtga aaggcatcct gtgcgggaag 360 420 cccqtqccqt catcttcttt qqtqaccaqq aqcatcccaa tqtcaccqag tttqctqtgg 480 qqcccctqcc aggqccctgc tacatgcgag cactgtcccc caggcctggg taccagtcct 540 cctqqqcatc qagqcccatc tccacagcag agtatgccct cctctaccac accctgcagg 600 aaqccaccaa gcccctgcat cagttcttcc tcaataccac aggcttctca ttccaagact 660 qccatgacag atgcctggcc ttcaccgatg tggcccccg gggtgtggct tCtggccagc 720 qccqcaqttq qcttatcata cagcgctatg tagaaggcta ctttctgcac cccactgggc 780 tgqaqctcct cgtggatcat gggagcacag atgctgggca ctgggccgtg gagcaggtgt ggtacaacgg gaagttctat gggagcccag aggaactggc tcggaagtat gcagatggag 840 900 aggtggacgt ggtggtcctg gaggacccgc tgcctggggg caaggggcat gacagcacag 960 aggagecgee cetettetee teccaeaage eeeggggga etteceeage eecatecatg 1020 tgagcggccc ccgcttggtc cagccccacg gccctcgctt caggctggag ggcaacgctg 1080 tgctctacqq cqqctqqaqc tttqccttcc ggctgcgctc ctcctccggg ctgcaggtcc tgaacgtgca cttcggcgga gagcgcattg cctatgaggt cagcgtgcaa gaggcagtgg 1140 cgctgtatgg aggacacaca cctgcaggca tgcagaccaa gtacctcgat gtcggctggg 1200 1260 qcctqqqcaq cqtcactcat qaqttagccc ccggcatcga ctgcccggag aCcgccacct tcctqqacac tttccactac tatqatgccq atqacccggt ccattatccc cgagccctct 1320 1380 gcctctttga aatgcccaca ggggtgcccc ttcggcggca ctttaattcc aactttaaag

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gagcctc
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<210> 25

<211> 1094

<212> DNA

<213> NM_000712.3| Homo sapiens biliverdin reductase A (BLVRA), mRNA

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| tgacctggct | ggtctccctc | tttggggagc | tttctcttgt | gtctgccact | ttggaagagc | 660 |
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| gaaaggaaga | tcagtatatg | aaaatgacag | tgtgtctgga | gacagagaag | aaaagtccac | 720 |
| tgtcatggat | tgaagaaaaa | ggacctggtc | taaaacgaaa | cagatattta | agctt c catt | 780 |
| tcaagtctgg | gtccttggag | aatgtgccaa | atgtaggagt | gaataagaac | atatttctga | 840 |
| aagatcaaaa | tatatttgtc | cagaaactct | tgggccagtt | ctctgagaag | gaactggctg | 900 |
| ctgaaaagaa | acgcatcctg | cactgcctgg | ggcttgcaga | agaaatccag | aaata t tgct | 960 |
| gttcaaggaa | gtaagaggag | gaggtgatgt | agcacttcca | agatggcacc | agcat t tggt | 1020 |
| tcttctcaag | agttgaccat | tatctctatt | cttaaaatta | aacatgttgg | ggaaa c aaga | 1080 |
| aaaaaaaaa | aaaa | | | | | 1094 |

<210> 26

<211> 5546

<212> DNA

<213> NM_000933.2| Homo sapiens phospholipase C, beta 4 (PLCB4), transcript variant 1, mRNA

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|------|---------------------|------------|------------|------------|------------|------------------------|
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| 180 | aagtt ccctc | tggcagaagg | tgaatttaac | ccaaacctta | ataatcatgg | agtcttgaat |
| 240 | tgttt g aacc | gaatcctttg | atacgaggag | tttttgacag | gaaggaacag | ctttttgcaa |
| 300 | gtgaa g gcaa | acatggagaa | cttctttctg | atgagtttgg | ttcaaagtgg | caactgcctc |
| 360 | ccata ccaaa | cggtcgggag | caacagtatt | gctccctcat | gtgctagaat | ggaaggacag |
| 420 | atctggaagg | tcagaaaatg | tgttggaaaa | ctcttgaagc | atcttggctg | ggatcccaaa |
| 480 | cctacatggt | attagtttta | tctagtgaac | gtggcacaga | tgtgtctgca | gcggatagtt |
| 540 | tcata cacaa | ctgagatcaa | ggtagaaggc | ctaagcaatg | ccagaagtaa | ggctgaaaat |
| 600 | tgaaattggc | aaacactgga | atgcctcaag | gtccaatgac | aacaacgtca | cttcagggcc |
| 660 | catttgcatc | attactagaa | agttaggagt | gtaaaattcc | aacacaaatg | atttatgacc |
| 720 | ccagt ggaaa | ttaggtcttc | actcaaggag | tctttcaagc | gaaaaggtga | gggaaaaaca |
| 780 | tgaca caaaa | ttctatgaac | ttatgaaaag | cagcattttc | attgagccca | gaatgatgaa |
| 840 | acaaa actga | atcaatggag | tttcaaaaaa | tagaagatct | cggacagata | gatttgtcct |
| 900 | atcct cgatt | catcaacgag | tctaaatgaa | tagtgagctt | gtagaccaat | ttatttaacg |
| 960 | ttgag atgta | atgcagatca | caaaagggca | tttatgatgc | ttatttccat | gaatgaaatt |
| 1020 | tttgcagata | agtgatgggt | ccttatatca | agaaaaaagg | gaagatttga | tgaacctgat |
| 1080 | accaa gaaat | ttagaacttt | cctagatcgt | ccccagtctt | gatgaaaacg | tctgatgtca |

| ggaccatcct | ctggctcact | acttcatcag | ttcttcccat | aacacttatc | tcactggcag | 1140 |
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| acagttcggc | gggaagtctt | cggtagaaat | gtacagacag | gttctcctgg | ctggttgcag | 1200 |
| atgtgttgaa | cttgactgct | gggatggaaa | aggtgaagac | caagaaccaa | taataactca | 1260 |
| tggaaaagca | atgtgtacag | atatcctttt | taaggatgta | attcaagcca | tcaaggaaac | 1320 |
| tgcatttgtc | acatcagaat | atcctgtaat | tctctccttt | gaaaatcact | gcagcaaata | 1380 |
| tcaacagtac | aagatgtcca | aatattgcga | agatctattt | ggggatctcc | tgttgaaaca | 1440 |
| agcacttgaa | tcacatccac | ttgaaccagg | cagggctttg | ccatccccca | atgacctcaa | 1500 |
| aagaaaaata | ctcataaaaa | acaagcggct | gaaacctgaa | gttgaaaaaa | aacagctgga | 1560 |
| agctttgaga | agcatgatgg | aagctggaga | atctgcctcc | ccagcaaaca | tcttagagga | 1620 |
| cgataatgaa | gaggagatcg | aaagtgctga | ccaagaggag | gaagctcacc | ccgaattcaa | 1680 |
| atttggaaat | gaactttctg | ctgatgactt | gggtcacaag | gaagctgttg | caaatagcgt | 1740 |
| caagaagggc | ctggtcactg | tagaagatga | gcaggcgtgg | atggcatctt | ataaatatgt | 1800 |
| aggtgctacc | actaatatcc | atccatattt | gtccacaatg | atcaactacg | cccagcctgt | 1860 |
| aaagtttcaa | ggtttccatg | tggcagaaga | acgcaatatt | cattataaca | tgtcttcttt | 1920 |
| taatgaatca | gtcggtcttg | gctacttgaa | gacacatgca | attgaatttg | tcaattataa | 1980 |
| caaacggcaa | atgagtcgca | tttaccccaa | gggaggccga | gtcgattcca | gtaattacat | 2040 |
| gcctcagatt | ttctggaacg | ctggctgcca | gatggtttca | ctgaactatc | aaaccccaga | 2100 |
| tttagcgatg | caattgaatc | agggaaaatt | tgagtataat | ggatcgtgcg | ggtaccttct | 2160 |
| caaaccagat | ttcatgaggc | ggcctgatcg | aacatttgac | cccttctctg | aaactcctgt | 2220 |
| tgatggtgtt | attgcagcca | cttgctcagt | gcaggttata | tcaggtcaat | tcttatcaga | 2280 |
| taagaaaatt | ggcacctacg | tagaggtgga | tatgtatggg | ttgcccactg | acaccatacg | 2340 |
| taaggaattc | cgaactcgca | tggttatgaa | taatggactc | aatccagttt | acaatgaaga | 2400 |
| gtcatttgta | tttcggaagg | tgatcctgcc | ggacctggct | gtcttgagaa | tagctgtgta | 2460 |
| tgatgataac | aacaagctga | ttggccagag | gatcctcccg | cttgatggcc | tccaagccgg | 2520 |
| atatcgacac | atttcccttc | gaaatgaggg | aaataaacca | ttatcactac | caacaatttt | 2580 |
| ctgcaatatt | gttcttaaaa | catatgtgcc | tgatggattt | ggagatatcg | tggatgcttt | 2640 |
| atcagatcca | aagaaatttc | tctcaattac | agaaaagaga | gcagaccaaa | tgagagctat | 2700 |
| gggcattgaa | actagtgaca | tagccgacgt | gcccagtgac | acttccaaaa | atgacaagaa | 2760 |
| aggaaaggcc | aacaccgcca | aagcaaatgt | gacccctcag | agtagctctg | agctcagacc | 2820 |
| aaccaccacg | gctgccctgg | cctctggtgt | ggaagccaag | aaaggtattg | aacttatccc | 2880 |
| tcaagtaagg | atagaagact | taaagcagat | gaaggcttac | ttgaagcatt | taaagaaaca | 2940 |
| gcagaaggag | ctaaattctt | taaagaagaa | acatgcaaag | gaacacagta | ccatgcagaa | 3000 |
| gttacactgc | acgcaagttg | acaaaattgt | ggcacagtat | gacaaagaga | agtcgactca | 3060 |

| aaaagaaaca gaaat | tcaaaa ttcagacgc | t gacatcagat | cacaaatcta | aggtcaaaga | 3180 |
|------------------|------------------|--------------|------------|------------|------|
| gattgtagca cagca | acacaa aggaatggt | c agaaatgatc | aatacccaca | gtgctgagga | 3240 |
| gcaagaaatc cgaga | acctgc acctcagcc | a gcagtgtgag | ctgctgaaaa | agctactcat | 3300 |
| caatgcccac gagca | agcaaa cccagcagc | t gaaactgtcc | catgacaggg | aaagcaagga | 3360 |
| aatgcgagca cacca | aggcta agatttcta | t ggaaaatagc | aaagccatca | gccaagataa | 3420 |
| atctatcaag aataa | aagcag aacgggaaa | g gcgagtcagg | gagttaaaca | gcagcaacac | 3480 |
| taaaaagttt ctgga | aagaaa gaaagagac | t tgccatgaag | cagtccaaag | aaatggatca | 3540 |
| gttgaaaaaa gtcca | agcttg aacatctag | a attcctagag | aaacagaatg | agcagctttt | 3600 |
| gaaatcctgt catgo | cagtgt cccaaacgc | a aggcgaagga | gatgcagcag | atggtgaaat | 3660 |
| tggaagccga gatg | gaccgc agaccagca | a cagtagtatg | aaactccaaa | atgcaaactg | 3720 |
| aagcagcaaa cccad | caaagc atcaaaaga | c tcactcacaa | acttctgaac | acaaactcca | 3780 |
| tggatgaaag ctgt | ttattt tgtttcctt | t atgtgtaaac | aagatgatat | ctgaaaccag | 3840 |
| agagacttgg aatg | tctgac tgacttcta | t ttaacagctt | gagtattgca | tttccttggc | 3900 |
| caaacaaaaa tagc | tacaaa tccacaaaa | a tttactattc | cagtaaggca | gagtccaacc | 3960 |
| attgataata caac | ttaaac atgtttgct | a taaaatacca | tcacaagtaa | atgagcttgg | 4020 |
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| acaaaaaaca gtgca | attagc aatttcata | g caagtgcatg | cactaggaaa | agaaaactct | 4140 |
| gtctacaagt ttat | tagcag aagtggtgg | t ctgctagaca | aataattttg | caaaattttt | 4200 |
| ctacatctaa gtta | cctcat cagtaagtg | c catgtctcta | ccatgccatc | agaggctaat | 4260 |
| ttcctgtaaa agtt | gtggaa attgttaga | a caatagaaaa | atagagcagt | gtatgtgtgc | 4320 |
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| gacattgctt tatag | ggaatt gtttctgca | g attccggata | ttataattca | caccataaag | 4440 |
| attgtgaagt ggtta | attggc aaacgtttg | t aaatgtgacc | atgtataaag | tatttatact | 4500 |
| ctttaattca cact | gttaga gagcaaaat | c atctaagtat | tgccacatga | caagattagt | 4560 |
| aaacaggaat acta | gaacta tgtttgcat | g atacacaagc | accaataaag | actaatccat | 4620 |
| acacagttaa ccta | atgcca aataaatac | t ggttaaataa | atgtatggca | cagaatataa | 4680 |
| tttgactatc aaga | ctttta gcataatga | a aaaccctctc | tctatatata | tatgtgtata | 4740 |
| tgaattatgt gggc | attctt gatacttca | a gttctagttt | gaaaaaaata | cataactaat | 4800 |
| ttaattttac acaa | aaatat ttatgcaga | t tttcagaatt | tcatatcagg | aaatgacctt | 4860 |
| tttatgtctg ttaa | atatca aaacaattt | g ctacagtgtt | aatctgcatg | gtctttaagc | 4920 |
| ctgctgtagt tgag | ttgcag acagtgcat | g aaaaagtatt | ccgctgggaa | ttgagccatg | 4980 |
| ccaccaaagc caag | aggagc gcatggaaa | c ccggtagtct | agaactaatc | agattactga | 5040 |
| ttttagggca cagc | accaga tgaattgtt | g tatatgcttg | taaaaattga | ttctgtgtgt | 5100 |

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tcctctgaac aaagcggaga aaatgatgat accatcaata ttgaaattaa acttccaact
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                                                                    5220
                                                                    5280
attctagcaa atgttttgcc tttttcatat acatgatatc atcgttattt tcaaaggggg
cttattaata ccctcagcat gtttttcacc caaatgatgc aaaacatgca gattctagtt
                                                                    5340
                                                                    5400
gacttcagtt gtaatagact tgtttttctc ctatttatga tttgaagtgg attctgtaaa
atatctcttg ttcttagttt ccttatctgt aaaacagtgg agttagacta catatctttt
                                                                    5460
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                                                                    5520
                                                                    5546
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<210> 27

<211> 2545

<212> DNA

<213> NM_002416.1| Homo sapiens chemokine (C-X-C motif) ligand 9 (CXCL9), mRNA

| | | | | | | 400 37 |
|------|------------|------------|------------|------------|------------|------------------------|
| 60 | tggtgttctt | tgaagaaaag | tctatcacta | ggaactccat | ggagtgactt | <400> 27 atccaataca |
| 120 | agtagtgaga | aaggaacccc | attggagtgc | gctggttctg | gcatcatctt | ttcctcttgg |
| 180 | atccttgaaa | tccacctaca | caagggacta | cagcaccaac | gttcctgcat | aagggtcgct |
| 240 | tgctacactg | ttgaaatcat | tgcgagaaaa | aagcccttcc | aatttgcccc | gaccttaaac |
| 300 | actgattaaa | atgtgaagga | gattcagcag | tctaaaccca | ttcaaacatg | aagaatggag |
| 360 | acatcaaaaa | atgggaaaaa | aagcaaaaga | ccaaaagaaa | aacaggtcag | aagtgggaga |
| 420 | tacataagag | aaaagaagac | cgttctcgtc | aaaatctcaa | tgaaagttcg | aagaaagttc |
| 480 | accgctatca | ttttaattat | aaatgttcta | tctgtgttaa | caataagtat | accacttcac |
| 540 | aatttaaaac | ttgactagaa | gcttattaat | taatacaaag | ggatggcata | ttccaaagga |
| 600 | acgttaagaa | aagaatccaa | agttgatttt | aaagttagaa | aattgtaact | attactctga |
| 660 | ttcatcatgc | ccagttgaat | ctaccaccca | ctttgttctt | ctatgattgt | ttgttaaagg |
| 720 | ccacatccca | gttcacccaa | ctacacagat | atacccatgt | gattttagca | ttaaggccat |
| 780 | ctccagagag | gtactgcagc | aggcttccac | gagcagccct | ctgcctggaa | ctcacaacag |
| 840 | ccaagcagtt | tgctggtgag | cctgttagca | aagtcctaag | acatgtcagc | tatctgaggc |
| 900 | ttgaatcagc | acctctgtat | gtggccatca | ccaagctgct | ctggacctca | tgaaattgag |
| 960 | gggtatcacc | gattgttatt | gattcatgct | gtgtctgaga | cacacacaat | ctacaggcct |
| 1020 | agccatgtga | tggctttgga | gcctcctttc | ggctttcaga | accagtgtgt | actggagatc |
| 1080 | tgcttcattc | tgttcccctt | tatttctttt | ctgaccactt | cccgctcagg | ttccatcttg |
| 1140 | gtgcacctgt | cttctctcca | cagtgccttt | taccacaatg | ttctccatcc | aagtcagctc |

| catatgctct | gatttatctg | agtcaactcc | tttctcatct | tgtccccaac | accccacaga | 1200 |
|------------|------------|------------|------------|------------|------------|------|
| agtgctttct | tctcccaatt | catcctcact | cagtccagct | tagttcaagt | cctgcctctt | 1260 |
| aaataaacct | ttttggacac | acaaattatc | ttaaaactcc | tgtttcactt | ggttcagtac | 1320 |
| cacatgggtg | aacactcaat | ggttaactaa | ttcttgggtg | tttatcctat | ctctccaacc | 1380 |
| agattgtcag | ctccttgagg | gcaagagcca | cagtatattt | ccctgtttct | tccacagtgc | 1440 |
| ctaataatac | tgtggaacta | ggttttaata | atttttaat | tgatgttgtt | atgggcagga | 1500 |
| tggcaaccag | accattgtct | cagagcaggt | gctggctctt | tcctggctac | tccatgttgg | 1560 |
| ctagcctctg | gtaacctctt | acttattatc | ttcaggacac | tcactacagg | gaccagggat | 1620 |
| gatgcaacat | ccttgtcttt | ttatgacagg | atgtttgctc | agcttctcca | acaataagaa | 1680 |
| gcacgtggta | aaacacttgc | ggatattctg | gactgttttt | aaaaaatata | cagtttaccg | 1740 |
| aaaatcatat | aatcttacaa | tgaaaaggac | tttatagatc | agccagtgac | caaccttttc | 1800 |
| ccaaccatac | aaaaattcct | tttcccgaag | gaaaagggct | ttctcaataa | gcctcagctt | 1860 |
| tctaagatct | aacaagatag | ccaccgagat | ccttatcgaa | actcatttta | ggcaaatatg | 1920 |
| agttttattg | tccgtttact | tgtttcagag | tttgtattgt | gattatcaat | taccacacca | 1980 |
| tctcccatga | agaaagggaa | cggtgaagta | ctaagcgcta | gaggaagcag | ccaagtcggt | 2040 |
| tagtggaagc | atgattggtg | cccagttagc | ctctgcagga | tgtggaaacc | tccttccagg | 2100 |
| ggaggttcag | tgaattgtgt | aggagaggtt | gtctgtggcc | agaatttaaa | cctatactca | 2160 |
| ctttcccaaa | ttgaatcact | gctcacactg | ctgatgattt | agagtgctgt | ccggtggaga | 2220 |
| tcccacccga | acgtcttatc | taatcatgaa | actccctagt | tccttcatgt | aacttccctg | 2280 |
| aaaaatctaa | gtgtttcata | aatttgagag | tctgtgaccc | acttaccttg | catctcacag | 2340 |
| gtagacagta | tataactaac | aaccaaagac | tacatattgt | cactgacaca | cacgttataa | 2400 |
| tcatttatca | tatatataca | tacatgcata | cactctcaaa | gcaaataatt | tttcacttca | 2460 |
| aaacagtatt | gacttgtata | ccttgtaatt | tgaaatattt | tctttgttaa | aatagaatgg | 2520 |
| tatcaataaa | tagaccatta | atcag | | | | 2545 |
| | | | | | | |

<210> 28

<211> 1144

<212> DNA

<213> NM_005859.2| Homo sapiens purine-rich element binding protein A (PURA), mRNA

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tggcggaccg agacagcggc agcgagcagg gtggtgcggc gctgggttcg ggcggctccc 120
tggggcaccc cggctcgggc tcaggctccg gcgggggcgg tggtggcggc gggggcggcg 180

| acaacsataa | caacaacaac | ggcggggccc | caggggggct | gcagcacgag | acgcaggagc | 240 |
|------------|------------|------------|------------|------------|------------|------|
| | | | | | | 700 |
| tggcctccaa | gcgggtggac | atccagaaca | agcgcttcta | cctggacgtg | aagcagaacg | 300 |
| ccaagggccg | cttcctgaag | atcgccgagg | tgggcgcggg | cggcaacaag | agccgcctta | 360 |
| ctctctccat | gtcagtggcc | gtggagttcc | gcgactacct | gggcgacttc | atcgagcact | 420 |
| acgcgcagct | gggccccagc | cagccgccgg | acctggccca | ggcgcaggac | gagccgcgcc | 480 |
| gggcgctcaa | aagcgagttc | ctggtgcgcg | agaaccgcaa | gtactacatg | gatctcaagg | 540 |
| agaaccagcg | cggccgcttc | ctgcgcatcc | gccagacggt | caaccggggg | cctggcctgg | 600 |
| gctccacgca | gggccagacc | attgcgctgc | ccgcgcaggg | gctcatcgag | ttccgtgacg | 660 |
| ctctggccaa | gctcatcgac | gactacggag | tggaggagga | gccggccgag | ctgcccgagg | 720 |
| gcacctcctt | gactgtggac | aacaagcgct | tcttcttcga | tgtgggctcc | aacaagtacg | 780 |
| gcgtgtttat | gcgagtgagc | gaggtgaagc | ccacctatcg | caactccatc | accgtgccct | 840 |
| acaaggtgtg | ggccaagttc | ggacacacct | tctgcaagta | ctcggaggag | atgaagaaga | 900 |
| ttcaagagaa | gcagagggag | aagcgggctg | cctgtgagca | gcttcaccag | cagcaacagc | 960 |
| agcagcagga | ggagaccgcc | gctgccaccc | tgctactgca | gggtgaggaa | gaaggggaag | 1020 |
| aagattgatc | aaactgaatg | aaacccccac | acacacacac | atgcatacac | acacacacac | 1080 |
| agccacacac | acagaaaata | tactgtaaag | aaagagagaa | aataaaaagt | taaaaagtta | 1140 |
| aaaa | | | | | | 1144 |
| | | | | | | |

<210> 29

<211> 1575

<212> DNA

 $<\!\!213\!\!>$ NM_014298.3| Homo sapiens quinolinate phosphoribosyltransferase (nicotinate-nucleotide pyrophosphorylase (carboxylating)) (QPRT), mRNA

| <400> 29 tcccaccccc | agcctggggc | ctctgggagc | cttggtcctg | agcagccaac | acaccagccc | 60 |
|------------------------|------------|------------|------------|------------|------------|-----|
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| ctggcagccc | tggtggacag | ctggctccga | gaggactgcc | cagggctcaa | ctacgcagcc | 180 |
| ttggtcagcg | gggcaggccc | ctcgcaggcg | gcgctgtggg | ccaaatcccc | tggggtactg | 240 |
| gcagggcagc | ctttcttcga | tgccatattt | acccaactca | actgccaagt | ctcctggttc | 300 |
| ctccccgagg | gatcgaagct | ggtgccggtg | gccagagtgg | ccgaggtccg | gggccctgcc | 360 |
| cactgcctgc | tgctggggga | acgggtggcc | ctcaacacgc | tggcccgctg | cagtggcatt | 420 |
| gccagtgctg | ccgccgctgc | agtggaggcc | gccagggggg | ccggctggac | tgggcacgtg | 480 |
| gcaggcacga | ggaagaccac | gccaggcttc | cggctggtgg | agaagtatgg | gctcctggtg | 540 |
| ggcggggccg | cctcgcaccg | ctacgacctg | ggagggctgg | tgatggtgaa | ggataaccat | 600 |

| gtggtggccg | ccggtggcgt | ggagaaggcg | gtgcgggcgg | ccagacaggc | ggctgacttc | 660 |
|------------|------------|------------|------------|------------|------------|------|
| gctctgaagg | tggaagtgga | atgcagcagc | ctgcaggagg | ccgtgcaggc | agctgaggct | 720 |
| ggtgccgacc | ttgtcctgct | ggacaacttc | aagccagagg | agctgcaccc | cacggccacc | 780 |
| gtgctgaagg | cccagttccc | gagtgtggct | gtggaagcca | gtgggggcat | caccctggac | 840 |
| aacctccccc | agttctgcgg | gccgcacata | gacgtcatct | ccatggggat | gctgacccag | 900 |
| gcggccccag | cccttgattt | ctccctcaag | ctgtttgcca | aagaggtggc | tccagtgccc | 960 |
| aaaatccact | agtcctaaac | cggaagagga | tgacaccggc | catgggttaa | cgtggctcct | 1020 |
| caggaccctc | tgggtcacac | atctttaggg | tcagtggcca | atggggcaca | tttggcacta | 1080 |
| gcttgagccc | aactctggct | ctgccacctg | ctgctcctgt | gacctgtcag | ggctgacttc | 1140 |
| acctctgctc | atctcagttt | cctaatctgt | aaaatgggtc | taataaagga | tcaaccacat | 1200 |
| ggggttctgc | ggtgataatg | agcacatagt | gaggggtcag | caaatgtcag | aagttacctg | 1260 |
| ggacagccgg | gcacgatggc | tcacacctgt | aatcccagca | ctttgggagg | ctgaggcggg | 1320 |
| aagatcactt | gagttcagga | gtttgagacc | agcctggcca | acatggtgaa | accccatctc | 1380 |
| taccaaaaat | agaagaatta | gctgggtgtg | gtggcacgcg | cctgtaatcc | cagctactta | 1440 |
| ggaggctgag | gcaggagaat | cgcttgaacc | caggaagtgg | aggttgcagt | gagctgatgg | 1500 |
| tgccactgca | ctccagcctg | ggtgatagag | cgagactctg | tctccaaaga | agaaaaaaaa | 1560 |
| aaaaaaaaa | aaaaa | | | | | 1575 |

<210> 30

<211> 768

<212> DNA

 $<\!\!213\!\!>$ NM_004585.2| Homo sapiens retinoic acid receptor responder (tazarotene induced) 3 (RARRES3), mRNA

| <400> 30 ccttcagcat | aaaagctgat | ccacaaacaa | gaggagcacc | agacctcctc | ttggcttcga | 60 |
|------------------------|------------|------------|------------|------------|------------|-----|
| gatggcttcg | ccacaccaag | agcccaaacc | tggagacctg | attgagattt | tccgccttgg | 120 |
| ctatgagcac | tgggccctgt | atataggaga | tggctacgtg | atccatctgg | ctcctccaag | 180 |
| tgagtacccc | ggggctggct | cctccagtgt | cttctcagtc | ctgagcaaca | gtgcagaggt | 240 |
| gaaacggggg | cgcctggaag | atgtggtggg | aggctgttgc | tatcgggtca | acaacagctt | 300 |
| ggaccatgag | taccaaccac | ggcccgtgga | ggtgatcatc | agttctgcga | aggagatggt | 360 |
| tggtcagaag | atgaagtaca | gtattgtgag | caggaactgt | gagcactttg | tcgcccagct | 420 |
| gagatatggc | aagtcccgct | gtaaacaggt | ggaaaaggcc | aaggttgaag | tcggtgtggc | 480 |
| cacggcgctt | ggaatcctgg | ttgttgctgg | atgctctttt | gcgattagga | gataccaaaa | 540 |
| aaaagcaaca | gcctgaagca | gccacaaaat | cctgtgttag | aagcagctgt | gggggtccca | 600 |

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660
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<211>
      696
<212>
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<213>
<400>
      31
                                                                    60
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                                                                   180
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ccaatgggct cagacctcc caccgcctgc tgcttttctt acaccgcgag gaagcttcct
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ttccaaacca aaagaagcaa gcaagtctgt gctgatccca gtgaatcctg ggtccaggag
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                                                                   420
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cacctgagcc cggatgcttc tccatgagac acatctcctc catactcagg actcctctcc
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                                                                   540
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| ttaaccgtcc | ttcccctagc | aaatttaaaa | acagaaagaa | aatgttgtac | cagttaccat | 2880 |
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| caaaaggagt | ggagcacagc | gtccggccca | gtgtgtttcc | ggttctgagt | cagggtgatc | 3120 |
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member 6 (TNFRSF6), transcript variant 1, mRNA

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| aatatcacca | aaaaatactt | aatagtccac | caaaaggcaa | gactgccctt | agaaattcta | 2220 |
| gcctggtttg | gagatactaa | ctgctctcag | agaaagtagc | tttgtgacat | gtcatgaacc | 2280 |
| catgtttgca | atcaaagatg | ataaaataga | ttcttatttt | tccccaccc | ccgaaaatgt | 2340 |
| tcaataatgt | cccatgtaaa | acctgctaca | aatggcagct | tatacatagc | aatggtaaaa | 2400 |
| tcatcatctg | gatttaggaa | ttgctcttgt | cataccccca | agtttctaag | atttaagatt | 2460 |
| ctccttacta | ctatcctacg | tttaaatatc | tttgaaagtt | tgtattaaat | gtgaatttta | 2520 |
| agaaataata | tttatatttc | tgtaaatgta | aactgtgaag | atagttataa | actgaagcag | 2580 |
| atacctggaa | ccacctaaag | aacttccatt | tatggaggat | ttttttgccc | cttgtgtttg | 2640 |
| gaattataaa | atataggtaa | aagtacgtaa | ttaaataatg | tttttggtaa | aaaaaaaaa | 2700 |
| aaaaaaaaa | aaaaaaaaaa | aaaaaaaaa | aaaaaaaaa | aaaaaaaaa | aaaaa | 2755 |

<210> 38

<211> 1600

<212> DNA

<213> NM_001953.2| Homo sapiens endothelial cell growth factor 1 (platelet-derived) (ECGF1), mRNA $\,$

| <400> 38 gccccgccgc | cggcagtgga | ccgctgtgcg | cgaaccctga | accctacggt | cccgacccgc | 60 |
|---------------------|--------------------|------------|------------|------------|------------|-----|
| gggcgaggcc | gggtacctgg | gctgggatcc | ggagcaagcg | ggcgagggca | gcgccctaag | 120 |
| caggcccgga | gcgatggcag | ccttgatgac | cccgggaacc | ggggccccac | ccgcgcctgg | 180 |
| tgacttctcc | ggggaaggga | gccagggact | tcccgaccct | tcgccagagc | ccaagcagct | 240 |
| cccggagctg | atccgcatga | agcgagacgg | aggccgcctg | agcgaagcgg | acatcagggg | 300 |
| cttcgtggcc | gctgtggtga | atgggagcgc | gcagggcgca | cagatcgggg | ccatgctgat | 360 |
| ggccatccga | cttcggggca | tggatctgga | ggagacctcg | gtgctgaccc | aggccctggc | 420 |
| tcagtcggga | cagcagctg g | agtggccaga | ggcctggcgc | cagcagcttg | tggacaagca | 480 |
| ttccacaggg | ggtgtgggtg | acaaggtcag | cctggtcctc | gcacctgccc | tggcggcatg | 540 |
| tggctgcaag | gtgccaatga | tcagcggacg | tggtctgggg | cacacaggag | gcaccttgga | 600 |
| taagctggag | tctattcctg | gattcaatgt | catccagagc | ccagagcaga | tgcaagtgct | 660 |
| gctggaccag | gcgggctgct | gtatcgtggg | tcagagtgag | cagctggttc | ctgcggacgg | 720 |
| aatcctatat | gcagccagag | atgtgacagc | caccgtggac | agcctgccac | tcatcacagc | 780 |
| ctccattctc | agtaagaaac | tcgtggaggg | gctgtccgct | ctggtggtgg | acgttaagtt | 840 |

| cggaggggcc | gccgtcttcc | ccaaccagga | gcaggcccgg | gagctggcaa | agacgctggt | 900 |
|------------|------------|------------|------------|------------|------------|------|
| tggcgtggga | gccagcctag | ggcttcgggt | cgcggcagcg | ctgaccgcca | tggacaagcc | 960 |
| cctgggtcgc | tgcgtgggcc | acgccctgga | ggtggaggag | gcgctgctct | gcatggacgg | 1020 |
| cgcaggcccg | ccagacttaa | gggacctggt | caccacgctc | gggggcgccc | tgctctggct | 1080 |
| cagcggacac | gcggggactc | aggctcaggg | cgctgcccgg | gtggccgcgg | cgctggacga | 1140 |
| cggctcggcc | cttggccgct | tcgagcggat | gctggcggcg | cagggcgtgg | atcccggtct | 1200 |
| ggcccgagcc | ctgtgctcgg | gaagtcccgc | agaacgccgg | cagctgctgc | ctcgcgcccg | 1260 |
| ggagcaggag | gagctgctgg | cgcccgcaga | tggcaccgtg | gagctggtcc | gggcgctgcc | 1320 |
| gctggcgctg | gtgctgcacg | agctcggggc | cgggcgcagc | cgcgctgggg | agccgctccg | 1380 |
| cctgggggtg | ggcgcagag⊂ | tgctggtcga | cgtgggtcag | aggctgcgcc | gtgggacccc | 1440 |
| ctggctccgc | gtgcaccggg | acggccccgc | gctcagcggc | ccgcagagcc | gcgccctgca | 1500 |
| ggaggcgctc | gtactctccg | accgcgcgcc | attcgccgcc | ccctcgccct | tcgcagagct | 1560 |
| cgttctgccg | ccgcagcaat | aaagctcctt | tgccgcgaaa | | | 1600 |
| | | | | | | |

<210> 39

<211> 931

<212> DNA

<213> NM_005138.1| Homo sapiens SCO cytochrome oxidase deficient homolog 2 (yeast) (SCO2), nuclear gene encoding mitochondrial protein, mRNA

| | | | | | | 400 00 | |
|-----|------------|------------|------------|------------|--------------------|------------------------|---|
| 60 | tgacgcgcgg | cttggtccac | tcctctcgtg | ggtcggcgct | gggagctgga | <400> 39 gcagagccca | ç |
| 120 | cagcttggca | cggagcccca | gctgctgact | gatccatgct | aggagcatca | cccgccgcg | C |
| 180 | aggccctgca | ctgggaggcc | ccctgggacc | ctccggtcct | cagctcaagc | caggctctct | (|
| 240 | agggccagcc | acaggtgggc | ccctgcagag | caaggcaggg | tggcttttgt | tctgaggtcc | t |
| 300 | ctggactcgg | ctgttcgggg | gatcacaggc | cccggctgct | gggcttcgaa | ccagggccct | (|
| 360 | agcgaacaga | cagcagcaaa | ggagaggctg | gggctgagaa | ctggccctga | tggggcctgg | 1 |
| 420 | acagaggccg | ctgctggatc | cgacttccac | tgggccaggg | caggcagctg | agccctgcgc | ć |
| 480 | gcttcactca | atgtactttg | gtgggtgctg | tccggggcca | aaggctgact | ggctcgctgc | ç |
| 540 | ggcagctgga | caggtggtgc | gaagctggtg | acgagctgga | atctgcccag | ctgccctgac | (|
| 600 | ccgagcggga | actgtggacc | tgtcttcatc | cagtgcagcc | ggtttgcctc | agcagagcct | ä |
| 660 | tgggtctgac | ccaagactgt | ggacttccac | gctacgtcca | gccatggccc | cgacgttgaa | (|
| 720 | acaatgccgg | cgcgtgtact | tcacagttac | cccaggctag | aaacaggtt g | cggctccacc | (|
| 780 | acctgctcaa | attgccatct | ggaccactcc | actacatcgt | gaggaccagg | ccccaaggat | (|
| 840 | agatctcaga | tcggctgagc | ccggagcaga | attactacgg | ctcttcacgg | ccctgacggc | (|

cagtgtgcgg cggcacatgg cggctttccg cagtgtcctg tcttgagcca ctgcagtctg 900 931 ggccccatca ttaaacgggc tgcgtttaaa a 40 <210> <211> 1216 <212> DNA NM_006419.1| Homo sapiens chemokine (C-X-C motif) ligand 13 (B-cell <213> chemoattractant) (CXCL13), mRNA <400> 40 60 ttcggcactt gggagaagat gtttgaaaaa actgactctg ctaatgagcc tggactcaga gctcaagtct gaactctacc tccagacaga atgaagttca tctcgacatc tctgcttctc 120 atgctgctgg tcagcagcct ctctccagtc caaggtgttc tggaggtcta ttacacaagc 180 ttgaggtgta gatgtgtcca agagagctca gtctttatcc ctagacgctt cattgatcga 240 300 attcaaatct tqccccqtqq qaatqqttqt ccaagaaaag aaatcatagt ctggaagaag aacaagtcaa ttgtgtgtgt ggaccctcaa gctgaatgga tacaaagaat gatggaagta 360 ttgagaaaaa gaagttcttc aactctacca gttccagtgt ttaagagaaa gattccctga 420 480 tgctgatatt tccactaaga acacctgcat tcttccctta tccctgctct ggattttagt 540 tttgtgctta gttaaatctt ttccagggag aaagaacttc cccatacaaa taaggcatga 600 ggactatgtg aaaaataacc ttgcaggagc tgatggggca aactcaagct tcttcactca cagcacceta tatacacttg gagtttgcat tettatteat cagggaggaa agtttetttg 660 aaaatagtta ttcagttata agtaatacag gattattttg attatatact tgttgtttaa 720 tgtttaaaat ttcttagaaa acaatggaat gagaatttaa gcctcaaatt tgaacatgtg 780 gcttgaatta agaagaaaat tatggcatat attaaaagca ggcttctatg aaagactcaa 840 aaagctgcct gggaggcaga tggaacttga gcctgtcaag aggcaaagga atccatgtag 900 tagatatcct ctgcttaaaa actcactacg gaggagaatt aagtcctact tttaaagaat 960 1020 ttctttataa aatttactgt ctaagattaa tagcattcga agatccccag acttcataga atactcaggg aaagcattta aagggtgatg tacacatgta tcctttcaca catttgcctt 1080 gacaaacttc tttcactcac atctttttca ctgacttttt ttgtgggggc ggggccgggg 1140 ggactctggt atctaattct ttaatgattc ctataaatct aatgacattc aataaagttg 1200 1216 agcaaacatt ttactt <210> 41 <211> 738 <212> DNA

<213> NM_006433.2| Homo sapiens granulysin (GNLY), transcript variant NKG5, mRNA

| <400> 41 gtatctgtgg taaacccagt gacacggggg agatgacata caaaaagggc aggacctgag | 60 |
|--|-----|
| aaagattaag ctgcaggctc cctgcccata aaacagggtg tgaaaggcat ctcagcggct | 120 |
| gccccaccat ggctacctgg gccctcctgc tccttgcagc catgctcctg ggcaacccag | 180 |
| gtctggtctt ctctcgtctg agccctgagt actacgacct ggcaagagcc cacctgcgtg | 240 |
| atgaggagaa atcctgcccg tgcctggccc aggagggccc ccagggtgac ctgttgacca | 300 |
| aaacacagga gctgggccgt gactacagga cctgtctgac gatagtccaa aaactgaaga | 360 |
| agatggtgga taagcccacc cagagaagtg tttccaatgc tgcgacccgg gtgtgtagga | 420 |
| cggggaggtc acgatggcgc gacgtctgca gaaatttcat gaggaggtat cagtctagag | 480 |
| ttacccaggg cctcgtggcc ggagaaactg cccagcagat ctgtgaggac ctcaggttgt | 540 |
| gtataccttc tacaggtccc ctctgagccc tctcaccttg tcctgtggaa gaagcacagg | 600 |
| ctcctgtcct cagatcccgg gaacctcagc aacctctgcc ggctcctcgc ttcctcgatc | 660 |
| cagaatccac tctccagtct ccctccctg actccctctg ctgtcctccc ctctcacgag | 720 |
| aataaagtgt caagcaag | 738 |

<210> 42

<211> 1579

<212> DNA

<213> NM_001767.2| Homo sapiens CD2 antigen (p50), sheep red blood cell receptor (CD2), mRNA

<400> 60 accaacccct aagatgagct ttccatgtaa atttgtagcc agcttccttc tgattttcaa tgtttcttcc aaaggtgcag tctccaaaga gattacgaat gccttggaaa cctggggtgc 120 180 cttgggtcag gacatcaact tggacattcc tagttttcaa atgagtgatg atattgacga 240 tataaaatgg gaaaaaactt cagacaagaa aaagattgca caattcagaa aagagaaaga 300 gactttcaag gaaaaagata catataagct atttaaaaat ggaactctga aaattaagca tctgaagacc gatgatcagg atatctacaa ggtatcaata tatgatacaa aaggaaaaaa 360 tqtqttqqaa aaaatatttg atttgaagat tcaagagagg gtctcaaaac caaagatctc 420 ctggacttgt atcaacacaa ccctgacctg tgaggtaatg aatggaactg accccgaatt 480 aaacctgtat caagatggga aacatctaaa actttctcag agggtcatca cacacaagtg 540 gaccaccagc ctgagtgcaa aattcaagtg cacagcaggg aacaaagtca gcaaggaatc 600 660 cagtgtcgag cctgtcagct gtccagagaa aggtctggac atctatctca tcattggcat

| atgtggagga | ggcagcctct | tgatggtctt | tgtggcactg | ctcgttttct | atatcaccaa | 720 |
|------------|------------|------------|------------|------------|------------|------|
| aaggaaaaaa | cagaggagtc | ggagaaatga | tgaggagctg | gagacaagag | cccacagagt | 780 |
| agctactgaa | gaaaggggcc | ggaagcccca | acaaattcca | gcttcaaccc | ctcagaatcc | 840 |
| agcaacttcc | caacatcctc | ctccaccacc | tggtcatcgt | tcccaggcac | ctagtcatcg | 900 |
| tccccgcct | cctggacacc | gtgttcagca | ccagcctcag | aagaggcctc | ctgctccgtc | 960 |
| gggcacacaa | gttcaccagc | agaaaggccc | gccctcccc | agacctcgag | ttcagccaaa | 1020 |
| acctccccat | ggggcagcag | aaaactcatt | gtccccttcc | tctaattaaa | aaagatagaa | 1080 |
| actgtctttt | tcaataaaaa | gcactgtgga | tttctgccct | cctgatgtgc | atatccgtac | 1140 |
| ttccatgagg | tgttttctgt | gtgcagaaca | ttgtcacctc | ctgaggctgt | gggccacagc | 1200 |
| cacctctgca | tcttcgaact | cagccatgtg | gtcaacatct | ggagtttttg | gtctcctcag | 1260 |
| agagctccat | cacaccagta | aggagaagca | atataagtgt | gattgcaaga | atggtagagg | 1320 |
| accgagcaca | gaaatcttag | agatttcttg | tccctctca | ggtcatgtgt | agatgcgata | 1380 |
| aatcaagtga | ttggtgtgcc | tgggtctcac | tacaagcagc | ctatctgctt | aagagactct | 1440 |
| ggagtttctt | atgtgccctg | gtggacactt | gcccaccatc | ctgtgagtaa | aagtgaaata | 1500 |
| aaagctttga | ctagaaaaaa | aaaaaaaaa | aaaaaaaaa | aaaaaaaaa | aaaaaaaaa | 1560 |
| aaaaaaaaa | aaaaaaaaa | | | | | 1579 |
| | | | | | | |

<210> 43

<211> 3738

<212> DNA

<213> NM_006275.4| Homo sapiens splicing factor, arginine/serine-rich 6 (SFRS6), mRNA

| 400 43 | | | | | | |
|---------------------|------------|------------|------------|------------|------------|-----|
| <400> 43 ctggcgcgcg | cgcgcgccat | tgtgtggctg | gactcggccg | cccctgtggt | gtgaggcgcg | 60 |
| tgttcgggct | cttgccgtcc | ccgcacccgc | accgcggtta | ctggcttgcg | gtccgccgtt | 120 |
| cgacaaccag | cccttgggtc | cccgcccgcc | acggacatgc | cgcgcgtcta | cataggacgc | 180 |
| ctgagctaca | acgtccggga | gaaggacatc | cagcgctttt | tcagtggcta | tggccgcctc | 240 |
| ctcgaagtag | acctcaaaaa | tgggtacggc | ttcgtggagt | tcgaggactc | ccgcgacgcc | 300 |
| gacgacgccg | tttacgagct | gaacggcaag | gagctctgcg | gcgagcgcgt | gatcgtagag | 360 |
| cacgcccggg | gcccgcgtcg | cgatcgcgac | ggctacagct | acggaagccg | cagtggtgga | 420 |
| ggtggataca | gcagtcggag | aacatctggc | agagacaaat | acggaccacc | tgttcgtaca | 480 |
| gaatacaggc | ttattgtaga | aaatctttct | agtcggtgca | gttggcaaga | tttaaaggat | 540 |
| tttatgcgac | aagcaggtga | agtaacctat | gcggatgccc | acaaggaacg | aacaaatgag | 600 |
| ggtgtaattg | agtttcgctc | ctactctgac | atgaagcgtg | ctttggacaa | actggatggc | 660 |

| acagaaataa | atggcagaaa | tattaggctt | attgaagata | agccacgcac | a.agccatagg | 720 |
|------------|------------|------------|------------|------------|---------------------|------|
| cgatcttact | ctggaagcag | atccaggtct | cgatctagaa | gacggtcacg | a agtaggagt | 780 |
| cgcaggagca | gccgcagtag | atctcgaagt | atctcaaaaa | gtcgctcccg | ttccaggtcg | 840 |
| cggagcaaag | gtcgatcacg | ttctcgatca | aaaggcagga | aatctagatc | a .aagagcaaa | 900 |
| tctaagccca | agtctgatcg | gggctcccat | tcacattctc | gaagcagatc | taaggatgag | 960 |
| tatgagaaat | ctcgaagcag | gtctcggtcc | cgatccccca | aagaaaatgg | a aagggtgat | 1020 |
| ataaagtcaa | aatccagatc | aaggagccag | tcccgttcca | attcgccgct | acctgttcca | 1080 |
| ccctcaaagg | cccgttctgt | gtcccctcca | ccaaaaagag | ctacttcaag | atcccgttct | 1140 |
| agatctcgct | caaagtcaag | atcaaggtcc | aggtcgagtt | ccagagatta | actcagaact | 1200 |
| ccttgtttgc | acattattat | ggaacacttt | cctacttagg | cagttactct | tccatgttta | 1260 |
| tacttggcct | cttctgcaag | aggaatctct | tgaaaacagg | ggcacacaga | aatttgattt | 1320 |
| gtggccaaat | tggatgaaaa | agatgaggct | ctaaggaaat | ggtggcatga | agaccctctc | 1380 |
| ccttctttgt | agaattaaga | taactttgat | tttatagctt | ttgagctaac | gtaacttttg | 1440 |
| taaagattaa | gctcatttag | tgttgtttt | tttttttt | tttttttt | t ttttagtat | 1500 |
| ttcagcagga | tctgctggca | gggtttttt | gttttatttg | tttgcttatt | tttaaattaa | 1560 |
| ctgttttgag | ctttgaatac | ttaaggcttt | agagggagaa | cccaattttc | a attatgttg | 1620 |
| gctttttata | aagcttgagt | tatgtaagat | ttaaataaaa | gtttgctacc | aagatgattg | 1680 |
| ccttattgaa | taggtcacta | ttaaattcct | ttaaatgttg | atatctgcca | t ttgtggaaa | 1740 |
| caacgtaaat | tctacttaag | tgtaaacaag | gcaagcctca | gaccagcaat | aaattactca | 1800 |
| gtttggataa | cattattttg | tgcagttaat | caaatttgcc | aaagtcttta | tctgcccctt | 1860 |
| taacaagttg | agtaaaaata | aaaggtattt | tttagtcaat | gtgttccatg | attttgctta | 1920 |
| aattaatact | tttaagtaat | ggaacttttt | tcaaaggcaa | atttaaacta | t ttaagaaat | 1980 |
| agctcctaat | acttgggatc | ttgtttagag | aatccacttt | ctggaagttc | tcagcataat | 2040 |
| tagtgttgag | agtggttcag | ttgtctttaa | tgtttgtcat | gtggaaatgg | aagtagcctc | 2100 |
| tttttgttct | gaaattgagt | ttattcaaag | tgtaaaagca | catactgcat | t ttctgctga | 2160 |
| aagatcatta | tgtttaacag | gcacttaatc | tcagtaaagt | cagttgccag | ttaagttcca | 2220 |
| cccagtagtc | agtccccttt | gtagttagtg | ggattatttg | ataattggtt | agatcatact | 2280 |
| tgtaaatttt | aatgctttgt | gtaattggtt | tgaaaaacag | tgaaatgggt | aaacgcaaaa | 2340 |
| cttttgtact | ttattacgag | taaagtgtaa | tgagtactgt | ggaaaccaaa | t ttgaatact | 2400 |
| gcaaatttgt | aggagttact | aggttagcaa | ttagtccata | catccataag | c ctgatgagt | 2460 |
| tgaaattgca | gtttgagaag | tgaattaacc | ttacatccct | ttgttcagat | accttaaaag | 2520 |
| ttactttatt | taaaagcatt | tattaatctt | agtctgaaat | caaaatatag | attaattggc | 2580 |
| tcagctttaa | tacctttcta | ggaggtgtca | caatgtaggg | taccaagggt | tggattgtga | 2640 |
| tggggcatgg | tcgtacactg | ctcattgtgc | cacaggtgtg | actggaaagc | atgatattct | 2700 |
| | | | | | | |

| | | | | | | | 2760 |
|-----|----------|------------|------------|------------|------------|------------|------|
| agg | gttggtt | tgtagattca | aataatccag | aaatatacct | aataagattg | agtgaaaaat | 2760 |
| ttg | agtcaaa | tatctagggc | attcacagag | tagctgtgag | ttcttggtaa | tgtgaaaaag | 2820 |
| gcc | ttgtttt | tcagaaattc | ctgggtttcc | tgttaaaaaa | tcttaaagcc | caaccttagg | 2880 |
| aat | atagtgc | cccaaaaggc | ggatgcttct | tccattatct | tattttcttt | gatactttat | 2940 |
| tta | attagat | gtttataaag | aaatgggttt | atttttccag | cataaacctc | agaatttaag | 3000 |
| gaa | agaaaat | gatgtctgtt | gttatagttc | attgttttgc | ctactcagca | gaagtgatga | 3060 |
| ctc | ttaaaaa | ttggctttga | ccaaagttct | cttgttttca | gggaaagaac | ataaaagctt | 3120 |
| ttt | gaactac | agccttttta | aaagagggat | gggaggatat | tacagtaaga | aattaggctt | 3180 |
| tct | aaaagta | tgaaacatcc | ttcaactggg | ctctcttgtt | aataggacat | catatggtaa | 3240 |
| tag | actggtt | tgactatatt | gttagctgcc | acagtaagca | ggtcattgta | taggtaaatg | 3300 |
| cct | gcaccca | taattttcta | gtaatagcca | cgaccaattt | attaacagtc | agggcctatc | 3360 |
| ctt | gcctgta | gttctcagtc | actggatgca | caaaatcact | gtgtaacatt | ggctcacttg | 3420 |
| gtg | gagcatag | ggttgactga | taaaatgttt | aattcccttg | ctagcttgtg | agaagaatga | 3480 |
| gtt | gatgaca | tgctccatac | cagtggctag | atggagtatt | aaggtggagc | agaaaagaag | 3540 |
| tga | agaacatc | ttgattcccc | tttcttttac | ttgatggtgt | ttatgaacat | gccgtagtgc | 3600 |
| ctt | tatggcc | agtttgagtc | ctgcctactt | tgacttttac | gttcccattc | ctgtgttacc | 3660 |
| aco | ttcctcc | cgatttgttc | acctattttg | tgctttaaat | ctcaataaaa | tacttactga | 3720 |
| gga | aaaaaaa | aaaaaaaa | | | | | 3738 |
| | | | | | | | |

<210> 44

<211> 2033

<212> DNA

<213> NM_003212.1| Homo sapiens teratocarcinoma-derived growth factor 1 (TDGF1), mRNA $\,$

| 400 | | | | | | | |
|--------------|----------------|------------|------------|------------|------------|------------|-----|
| <400 ggag | > 44 aatccc | cggaaaggct | gagtctccag | ctcaaggtca | aaacgtccaa | ggccgaaagc | 60 |
| cctc | cagttt | cccctggacg | ccttgctcct | gcttctgcta | cgaccttctg | gggaaaacga | 120 |
| attt | ctcatt | ttcttcttaa | attgccattt | tcgctttagg | agatgaatgt | tttcctttgg | 180 |
| ctgt | tttggc | aatgactctg | aattaaagcg | atgctaacgc | ctcttttccc | cctaattgtt | 240 |
| aaaa | gctatg | gactgcagga | agatggcccg | cttctcttac | agtgtgattt | ggatcatggc | 300 |
| catt | tctaaa | gtctttgaac | tgggattagt | tgccgggctg | ggccatcagg | aatttgctcg | 360 |
| tcca | tctcgg | ggatacctgg | ccttcagaga | tgacagcatt | tggccccagg | aggagcctgc | 420 |
| aatt | cggcct | cggtcttccc | agcgtgtgcc | gcccatgggg | atacagcaca | gtaaggagct | 480 |
| aaac | agaaco | tgctgcctga | atgggggaac | ctgcatgctg | ggtcctttt | gtgcctgccc | 540 |

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| ccgctgcttt | cctcaggcat | ttctacccgg | ctgtgatggc | cttgtgatgg | atgagcacct | 720 |
| cgtggcttcc | aggactccag | aactaccacc | gtctgcacgt | actaccactt | ttatgctagt | 780 |
| tggcatctgc | ctttctatac | aaagctacta | ttaatcgaca | ttgacctatt | tccagaaata | 840 |
| caattttaga | tatcatgcaa | atttcatgac | cagtaaaggc | tgctgctaca | atgtcctaac | 900 |
| tgaaagatga | tcatttgtag | ttgccttaaa | ataatgaata | caatttccaa | aatggtctct | 960 |
| aacatttcct | tacagaacta | cttcttactt | ctttgccctg | ccctctccca | aaaaactact | 1 O 20 |
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| gttcagatta | ttggagacta | attctaatgt | ggaccttaga | atacagtttt | gagtagagtt | 1500 |
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| tggtggtaga | gaagcaagta | aaaaggctaa | atggaagggc | aagtttccat | catctataga | 1740 |
| aagctatata | agacaagaac | tccccttttt | ttcccaaagg | cattataaaa | agaatgaagc | 1800 |
| ctccttagaa | aaaaaattat | acctcaatgt | ccccaacaag | attgcttaat | aaattgtgtt | 1860 |
| tcctccaagc | tattcaattc | ttttaactgt | tgtagaagac | aaaatgttca | caatatattt | 1920 |
| agttgtaaac | caagtgatca | aactacatat | tgtaaagccc | atttttaaaa | tacattgtat | 1980 |
| atatgtgtat | gcacagtaaa | aatggaaact | atattgacct | aaaaaaaaa | aaa | 2 0 33 |
| .210. 45 | | | | | | |
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| <712> NW_(| 70383T'T H | omo saptens | metariotiii | опен ти (м | I III , IIINNA | |
| رام مرمد مارام مارام | | | | | | |
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<211> 3052

<212> DNA

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| caaaataccc | ccaacatacc | agatccgctt | cctgccccgc | tgaaggggct | gagggaaggg | 1500 |
| ggtcaaagga | ttccagggtc | attcagtgtc | cccgcctctg | tagacaatgg | ctctgactcc | 1560 |
| ccgcaacttc | ctgcctctga | gagacctgct | acaagccagc | ttccttcccc | tccatggcac | 1620 |
| cagttgtctg | aggtcacatt | gcaagtgagt | gcaggagtga | gattatcgaa | aattataata | 1680 |
| tacaaaatca | tatatata | tatgttcttg | ttttttgaga | cagagtctca | cactgttgcc | 1740 |
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| caactctcct | gcctcagcct | ccctagtagc | tgggattaca | ggcatgcact | accacgcttg | 1860 |
| gctaattttt | gtatttttag | tagagatggg | gtttcactgt | gtaggccagg | ctggtctcga | 1920 |
| actcctgaac | tcaagtgatt | cacccacctt | agcctcccaa | agtgctggga | ttacaggcgt | 1980 |
| gagtcaccgt | gcccagccat | gtatatatat | aattttaaaa | attaagctga | aattcacata | 2040 |
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<210> 47

<211> 1645

<212> DNA

 $<\!\!213\!\!>$ NM_003811.2| Homo sapiens tumor necrosis factor (ligand) superfamily, member 9 (TNFSF9), mRNA

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| ccgcgccccg | cgctcgcgcc | tgccgcgtac | tgccttgggc | cctggtcgcg | gggctgctgc | 120 |
| tgctgctgct | gctcgctgcc | gcctgcgccg | tcttcctcgc | ctgcccctgg | gccgtgtccg | 180 |
| gggctcgcgc | ctcgcccggc | tccgcggcca | gcccgagact | ccgcgagggt | cccgagcttt | 240 |
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| attctgagcc | tgagctcaga | taatatatta | tatatattat | atatatatat | atatttctat | 1020 |
| ttaaagagga | tcctgagttt | gtgaatggac | ttttttagag | gagttgtttt | 999999999 | 1080 |
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| cctcccaagc | aactgggatt | catcctttct | attaattcat | tgtacttatt | tgcctatttg | 1200 |
| tgtgtattga | gcatctgtaa | tgtgccagca | ttgtgcccag | gctagggggc | tatagaaaca | 1260 |
| tctagaaata | gactgaaaga | aaatctgagt | tatggtaata | cgtgaggaat | ttaaagactc | 1320 |
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| acggtattgc | tatgttgcca | aggttgttta | catgccagta | caatttataa | taaacactca | 1620 |
| tttttcctca | aaaaaaaaaa | aaaaa | | | | 1645 |

62

<210> 48 <211> 6640

<212> DNA

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| aagatagata | tgattcgaaa | aagactgcag | aacttcagct | atgaccagag | ggaaatgata | 1860 |
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| ctaaatccag | agggggatgt | caactctgcc | aaagtctgtg | cccacataac | aaatattcca | 1920 |
| ttcagcatta | caaagatgga | tgttcttcag | ttcctagaag | gaatcccagt | ggatgaaaat | 1980 |
| gctgtacatg | ttcttgttga | taacaatggg | caaggtctag | gacaggcatt | ggttcagttt | 2040 |
| aaaaatgaag | atgatgcacg | taagtctgaa | cgcttacacc | gtaaaaaact | taatgggaga | 2100 |
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| aaagtcttca | tacagagtac | aagtgcagcc | gccagaggag | aaaattgaga | ttcttgaccc | 4260 |
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| ccttgtgatg | aaaaacttgg | ccttacaggg | ctctagttat | ggttactttc | cctagtcaat | 4560 |
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| gagagatgca | gacaaagtaa | acaatgaagg | ttgttttata | aaggtgatga | ccattataga | 720 |
| gtcagaaatg | ggagtcgttg | caggaatttc | ctttggagtt | gcttgcttcc | aactgattgg | 780 |
| aatctttctc | gcctactgcc | tctctcgtgc | cataacaaat | aaccagtatg | agatagtgta | 840 |
| acccaatgta | tctgtgggcc | tattcctctc | tacctttaag | gacatttagg | gtccccctg | 900 |
| tgaattagaa | agttgcttgg | ctggagaact | gacaacacta | cttactgata | gaccaaaaaa | 960 |
| ctacaccagt | aggttgattc | aatcaagatg | tatgtagacc | taaaactaca | ccaataggct | 1020 |
| gattcaatca | agatccgtgc | tcgcagtggg | ctgattcaat | caagatgtat | gtttgctatg | 1080 |
| ttctaagtcc | accttctatc | ccattcatgt | tagatcgttg | aaaccctgta | tccctctgaa | 1140 |
| acactggaag | agctagtaaa | ttgtaaatga | agtaatactg | tgttcctctt | gactgttatt | 1200 |
| tttcttagta | gggggccttt | ggaaggcact | gtgaatttgc | tattttgatg | tagtgttaca | 1260 |
| agatggaaaa | ttgattcctc | tgactttgct | attgatgtag | tgtgatagaa | aattcacccc | 1320 |
| tctgaactgg | ctccttccca | gtcaaggtta | tctggtttga | ttgtataatt | tgcaccaaga | 1380 |
| agttaaaatg | ttttatgact | ctctgttctg | ctgacaggca | gagagtcaca | ttgtgtaatt | 1440 |
| taatttcagt | cagtcaatag | atggcatccc | tcatcagggt | tgccagatgg | tgataacagt | 1500 |
| gtaaggcctt | gggtctaagg | catccacgac | tggaagggac | tactgatgtt | ctgtgataca | 1560 |
| tcaggtttca | gcacacaact | tacatttctt | tgcctccaaa | ttgaggcatt | tattatgatg | 1620 |
| ttcatacttt | ccctcttgtt | tgaaagtttc | taattattaa | atggtgtcgg | aattgttgta | 1680 |
| ttttccttag | gaattcagtg | gaacttatct | tcattaaatt | tagctggtac | caggttgata | 1740 |
| tgacttgtca | atattatggt | caactttaag | tcttagtttt | cgtttgtgcc | tttgattaat | 1800 |
| aagtataact | cttatacaat | aaatactgct | ttcctctaaa | aagatcgtgt | ttaaattaac | 1860 |
| ttgtagaaaa | tctgctggaa | tggttgttgt | tttccactga | gaaagctaag | ccctacattt | 1920 |
| ctattcagag | tactgtttt | agatgtgaaa | tataagcctg | cggccttaac | tctgtattaa | 1980 |
| aaaaaatgtt | tttgtttaaa | aaaaactgtt | cccataggtg | cagcaaacca | ccatggcaca | 2040 |
| tgtataccta | tgtaacaaac | ctgcacatt | | | | 2069 |
| | | | | | | |

<210> 59

<211> 2402

<212> DNA

 $<\!\!213\!\!>\!$ NM_021200.1| Homo sapiens pleckstrin homology domain containing, family B (evectins) member 1 (PLEKHB1), mRNA

| <400> 59 aagagaggaa | ggcttaaaga | gccagactgc | gcagccagga | ctggggtatg | ggcgctgtcc | 60 |
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| tcctgtcctg | ccccgcaac | ctcgccccga | ttccactccg | ggaacctcgg | cgatgctgag | 180 |
| ccaagaccac | ttctgaatca | gggatgactt | gtctagtgaa | cgtagggtca | gagccatcag | 240 |
| ttggaaaggc | tgggaggagc | ctggagaaag | aggcgacctt | ccttgggatc | tgtgcgctcc | 300 |
| ctccttgcct | cccctccag | cctcccactt | ggtagcacct | tcctgatccc | cttatctcta | 360 |
| aggcgctcag | ggaaatgccc | cgctgcggga | gccttctggg | aaatgctgcc | ctggccaccc | 420 |
| aggaaccatg | agccctgcag | ccccggtccc | gcctgactcc | gctctggaaa | gtccttttga | 480 |
| agaaatggcc | ctggtgaggg | gcggctggct | gtggagacag | agctccatcc | tccgccgctg | 540 |
| gaagcggaac | tggtttgccc | tgtggctgga | cgggaccctg | ggatactacc | acgatgagac | 600 |
| agcgcaggac | gaggaggacc | gtgtgctcat | ccacttcaat | gtccgtgaca | taaagatcgg | 660 |
| cccagagtgc | catgatgtgc | agcccccaga | gggccggagc | cgagatggcc | tgctgactgt | 720 |
| gaacctacgg | gaaggcggcc | gcctgcacct | ctgtgcggag | accaaggatg | atgccctagc | 780 |
| atggaagaca | gcactgctgg | aggcaaactc | caccccggcc | ccagctggag | ccaccgtccc | 840 |
| tcccaggagc | cgccgggttt | gctccaaggt | caggtgtgtg | acccgctcgt | ggagcccctg | 900 |
| taaggttgag | aggcggatct | gggtgcgcgt | ctacagcccg | taccaagact | actacgaggt | 960 |
| ggtgccccc | aatgcacacg | aggccacgta | tgtccgcagc | tactacggac | cgccctacgc | 1020 |
| aggccctggc | gtgacgcacg | tgatagtgcg | ggaggatccc | tgctacagcg | ccggcgcccc | 1080 |
| tctggccatg | ggcatgcttg | cgggagccgc | cactggggcg | gcgctgggct | cgctcatgtg | 1140 |
| gtcgccctgc | tggttctgag | ccctgggact | cggagcactg | acccctgcgc | ttggattgct | 1200 |
| agactcctct | tcctcctgga | ccccatcctc | taccatccaa | gccctgtccc | actttggccc | 1260 |
| tatcctctcc | attagctcct | tccgggtttg | gaccattccc | cccactccct | acccttaatc | 1320 |
| cccacatggg | aagaagctat | catcacaggt | acaaacatcg | cttgaagtct | tcacatctac | 1380 |
| cactagacac | ccccaaaatc | tgttatagac | atttatggat | acatttcctc | taaacacaac | 1440 |
| agggcacagc | aaatacgact | tcatttggct | tcgagttccc | caggcgctgt | agacacaaca | 1500 |
| tgaatcgggc | tctctgctct | ctccttaggg | agctcgagtc | ctggtgggga | gaacaggagt | 1560 |
| aaacaaggac | ttgacaaagc | tgaagagtta | tcagtccttt | gacaaggaca | ggtggggcag | 1620 |
| ggagcaagac | aggtaggctg | gaagaacagt | tattggcaag | tatgcagagc | cgtgaacgtc | 1680 |
| atggcatgtc | caaggaatta | aatgggagtt | catttgggct | ggggtggagg | ctgggatcag | 1740 |
| accgtggtgg | gccttcaagc | taaggagctt | cctaggtgaa | aggggagatg | tgagccttct | 1800 |
| ctggagggaa | gtttcatgat | tgcatctata | atgaatatat | tgcctgtttt | gtgaatactg | 1860 |
| acacatgtcc | atacctaaaa | cactcctgag | ttaagtccca | tccttcccac | aaacagcttc | 1920 |
| ctggctggta | cccatgataa | caattgagct | gaacctgggg | acccctggtt | ggggaacagg | 1980 |

| tgagttctat | ttgagacttc | cagccctaga | aagctgcctc | cgtccagaaa | tgcctctcac | 2040 |
|------------|------------|------------|------------|------------|------------|------|
| accaggagct | cggccctctc | tttatagctg | tgactgtcac | cctctcaggc | tttgtctcat | 2100 |
| ccttcattct | gaataagatg | gcagtgttct | cctctggggc | ctgatccacc | tctacaccag | 2160 |
| cccaggaagc | cccatctgtg | cctgccctca | ggtggtccac | cagtctcccc | ctttggttcc | 2220 |
| cttccagtct | cttccccctt | tctatcccaa | tcaccaatag | aaatgctaac | atccctgcct | 2280 |
| ggtagccaga | ctagcccact | aaagctcccc | tgtaaatggg | ggctccatta | gttctgctgc | 2340 |
| cgagactaat | aaagatttgg | ttggttctag | cagtaaaaaa | aaaaaaaaa | aaaaaaaaaa | 2400 |
| aa | | | | | | 2402 |

<210> 60

<211> 2856

<212> DNA

 $<\!\!213\!\!>$ NM_003661.2| Homo sapiens apolipoprotein L, 1 (APOL1), transcript variant 1, mRNA

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|------|------------|------------|------------|------------|------------|---------------------|
| 120 | ttgctcagtc | cagctggatc | cagctcagaa | gggatccaca | aactggaggt | acagacgcat |
| 180 | gctgctttgc | catggaggga | cctgcagcga | tggaggaggc | gaagattcct | tctgccaggg |
| 240 | ggagtgaggg | ccttggtgtg | gtgcactttt | atctggatga | tgtcctctgc | tgagagtctc |
| 300 | actggagatc | tgggacagat | acgttccaag | gtgcaacaaa | tggagcgagg | cagaggaagc |
| 360 | agcagtatct | ggacccagag | ctggcaccat | gactgggctg | gcccctcggt | ctcaaagtaa |
| 420 | ctgctactcc | cacacagaat | aaaaagtgag | tatttcaagg | tgccattaag | ttattgagga |
| 480 | cccaggaatg | tgctgaactg | tcgtggctgc | tggaacggat | taatgaggcc | tgctgactga |
| 540 | atgaaagaca | acaaatgatc | accttgcaag | gctctggaca | gctccgtaaa | aggcagatga |
| 600 | tttcctcggt | tctgaaagag | gaaactggtt | cagcagtaca | cgataaaggc | aaaactggca |
| 660 | ggggttcaga | ccttgcagat | ggctccgtgc | aacataagaa | gcttgaggat | tgaaaagtga |
| 720 | atttcctctg | ctctctcagc | tggtgtctgg | atcgccaatg | aggcaccacc | aggtccacaa |
| 780 | agccttgtac | agagggaggc | cacccttcac | atgggtctgg | cctcgtcggc | gcatcctgac |
| 840 | accagcagta | gaccgggatt | cagccgcttt | ttgggaatca | tgggatggag | tcttggaacc |
| 900 | gtcatcaaaa | ccacgacctg | aagcccaagc | tggtggacac | cggaaagaag | ccatggacta |
| 960 | aactttcttt | gaacatatcc | ttttgggtga | gtgagggagt | attgaaggag | gccttgacaa |
| 1020 | cgtgccctca | gaaggacatc | gaggcattgg | caactcacac | caatacttac | ccttagctgg |
| 1080 | ccccgggtca | agcctcacgc | cgcatgcctc | cagtcagtac | agccaatctt | gacgagccag |
| 1140 | cccagcatcc | ggttaatgaa | aggtggagag | agcggtgaac | ctcagctgaa | ctgagccaat |

| tggaaatgag | cagaggagtc | aagctcacgg | atgtggcccc | tgtaagcttc | tttcttgtgc | 1200 |
|------------|------------|------------|------------|------------|------------|------|
| tggatgtagt | ctacctcgtg | tacgaatcaa | agcacttaca | tgagggggca | aagtcagaga | 1260 |
| cagctgagga | gctgaagaag | gtggctcagg | agctggagga | gaagctaaac | attctcaaca | 1320 |
| ataattataa | gattctgcag | gcggaccaag | aactgtgacc | acagggcagg | gcagccacca | 1380 |
| ggagagatat | gcctggcagg | ggccaggaca | aaatgcaaac | tttttttt | ttctgagaca | 1440 |
| gagtcttgct | ctgtcgccaa | gttggagtgc | aatggtgcga | tctcagctca | ctgcaagctc | 1500 |
| tgcctcccgt | gttcaagcga | ttctcctgcc | ttggcctccc | aagtagctgg | gactacaggc | 1560 |
| gcctaccacc | atgcccagct | aatttttgta | tttttaatag | agatggggtt | tcaccatgtt | 1620 |
| ggccaggatg | gtctcgatct | cctgacctct | tgatctgccc | accttggcct | cccaaagtgc | 1680 |
| tgggattaca | ggcgtgagcc | atcgcttttg | acccaaatgc | aaacatttta | ttagggggat | 1740 |
| aaagagggtg | aggtaaagtt | tatggaactg | agtgttaggg | actttggcat | ttccatagct | 1800 |
| gagcacagca | ggggaggggt | taatgcagat | ggcagtgcag | caaggagaag | gcaggaacat | 1860 |
| tggagcctgc | aataagggaa | aaatgggaac | tggagagtgt | ggggaatggg | aagaagcagt | 1920 |
| ttactttaga | ctaaagaata | tattgggggg | ccgggtgtag | tggctcatgc | ctgtaatccg | 1980 |
| agcactttgg | gaggccaagg | cgggcggatc | acgaggtcag | gagatcgaga | ccatcctggc | 2040 |
| taacacagtg | aaaccccgtc | tctactaaaa | atacaaaaaa | ttagccgggc | atggtggcgg | 2100 |
| gcgcctgtag | ttccagctaa | ctgggcggct | gaggcaggag | aatggcgtga | acctgggagg | 2160 |
| tggagcttgc | agtgagccga | gatatcgcca | ctgcactcca | gcctgggtga | cagagcgaga | 2220 |
| ctccatctca | aaaaaaaaa | aaaaaagaat | atattgacgg | aagaatagag | aggaggcttg | 2280 |
| aaggaaccag | caatgagaag | gccaggaaaa | gaaagagctg | aaaatggaga | aagcccaaga | 2340 |
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| atggtcattg | gggtggttgt | catgtgatgg | gtcccctcca | ggttactaaa | gggtgcatgt | 2580 |
| cccctgcttg | aacactgaag | ggcaggtggt | gggccatggc | catggtcccc | agctgaggag | 2640 |
| caggtgtccc | tgagaaccca | aacttcccag | agagtatgtg | agaaccaacc | aatgaaaaca | 2700 |
| gtcccatcgc | tcttacccgg | taagtaaaca | gtcagaaaat | tagcatgaaa | gcagtttagc | 2760 |
| attgggagga | agctcagatc | tctagagctg | tcttgtcgcc | gcccaggatt | gacctgtgtg | 2820 |
| taagtcccaa | taaactcacc | tactcatcaa | gctgga | | | 2856 |
| | | | | | | |

<210> 61

<211> 1655

<212> DNA

 $\ensuremath{^{<\!213>}}$ NM_002164.3| Homo sapiens indoleamine-pyrrole 2,3 dioxygenase (INDO), mRNA

| .400. C1 | | | | | | |
|------------------------|------------|------------|------------|------------|------------|------|
| <400> 61 aatttctcac | tgcccctgtg | ataaactgtg | gtcactggct | gtggcagcaa | ctattataag | 60 |
| atgctctgaa | aactcttcag | acactgaggg | gcaccagagg | agcagactac | aagaatggca | 120 |
| cacgctatgg | aaaactcctg | gacaatcagt | aaagagtacc | atattgatga | agaagtgggc | 180 |
| tttgctctgc | caaatccaca | ggaaaatcta | cctgattttt | ataatgactg | gatgttcatt | 240 |
| gctaaacatc | tgcctgatct | catagagtct | ggccagcttc | gagaaagagt | tgagaagtta | 300 |
| aacatgctca | gcattgatca | tctcacagac | cacaagtcac | agcgccttgc | acgtctagtt | 360 |
| ctgggatgca | tcaccatggc | atatgtgtgg | ggcaaaggtc | atggagatgt | ccgtaaggtc | 420 |
| ttgccaagaa | atattgctgt | tccttactgc | caactctcca | agaaactgga | actgcctcct | 480 |
| attttggttt | atgcagactg | tgtcttggca | aactggaaga | aaaaggatcc | taataagccc | 540 |
| ctgacttatg | agaacatgga | cgttttgttc | tcatttcgtg | atggagactg | cagtaaagga | 600 |
| ttcttcctgg | tctctctatt | ggtggaaata | gcagctgctt | ctgcaatcaa | agtaattcct | 660 |
| actgtattca | aggcaatgca | aatgcaagaa | cgggacactt | tgctaaaggc | gctgttggaa | 720 |
| atagcttctt | gcttggagaa | agcccttcaa | gtgtttcacc | aaatccacga | tcatgtgaac | 780 |
| ccaaaagcat | ttttcagtgt | tcttcgcata | tatttgtctg | gctggaaagg | caacccccag | 840 |
| ctatcagacg | gtctggtgta | tgaagggttc | tgggaagacc | caaaggagtt | tgcagggggc | 900 |
| agtgcaggcc | aaagcagcgt | ctttcagtgc | tttgacgtcc | tgctgggcat | ccagcagact | 960 |
| gctggtggag | gacatgctgc | tcagttcctc | caggacatga | gaagatatat | gccaccagct | 1020 |
| cacaggaact | tcctgtgctc | attagagtca | aatccctcag | tccgtgagtt | tgtcctttca | 1080 |
| aaaggtgatg | ctggcctgcg | ggaagcttat | gacgcctgtg | tgaaagctct | ggtctccctg | 1140 |
| aggagctacc | atctgcaaat | cgtgactaag | tacatcctga | ttcctgcaag | ccagcagcca | 1200 |
| aaggagaata | agacctctga | agacccttca | aaactggaag | ccaaaggaac | tggaggcact | 1260 |
| gatttaatga | atttcctgaa | gactgtaaga | agtacaactg | agaaatccct | tttgaaggaa | 1320 |
| ggttaatgta | acccaacaag | agcacatttt | atcatagcag | agacatctgt | atgcattcct | 1380 |
| gtcattaccc | attgtaacag | agccacaaac | taatactatg | caatgtttta | ccaataatgc | 1440 |
| aatacaaaag | acctcaaaat | acctgtgcat | ttcttgtagg | aaaacaacaa | aaggtaatta | 1500 |
| tgtgtaatta | tactagaagt | tttgtaatct | gtatcttatc | attggaataa | aatgacattc | 1560 |
| aataaataaa | aaaaaaaaa | aaaaaaaaa | aaaaaaaaa | aaaaaaaaa | aaaaaaaaa | 1620 |
| aaaaaaaaa | aaaaaaaaa | aaaaaaaaa | aaaaa | | | 1655 |
| | | | | | | |

<210> 62

<211> 2242

<212> DNA

 $<\!\!213\!\!>$ NM_021784.3| Homo sapiens forkhead box A2 (FOXA2), transcript variant 1, mRNA

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| aaaagagggt | gggggtgggg | ggtgattgct | ggtcgtttgt | tgtggctgtt | aaattttaaa | 180 |
| ctgccatgca | ctcggcttcc | agtatgctgg | gagcggtgaa | gatggaaggg | cacgagccgt | 240 |
| ccgactggag | cagctactat | gcagagcccg | agggctactc | ctccgtgagc | aacatgaacg | 300 |
| ccggcctggg | gatgaacggc | atgaacacgt | acatgagcat | gtcggcggcc | gccatgggca | 360 |
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| tcggggggca | ggcggccggg | gccatgggcg | gcctggcccc | ctacgccaac | atgaactcca | 600 |
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| ggaccctgca | ccctgactcg | ggcaacatgt | tcgagaacgg | ctgctacctg | cgccgccaga | 960 |
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| gcaagaaggc | ggccgccgga | gcccaggcct | cacaggctca | actcggggag | gccgccgggc | 1080 |
| cggcctccga | gactccggcg | ggcaccgagt | cgcctcactc | gagcgcctcc | ccgtgccagg | 1140 |
| agcacaagcg | agggggcctg | ggagagctga | aggggacgcc | ggctgcggcg | ctgagccccc | 1200 |
| cagagccggc | gccctctccc | gggcagcagc | agcaggccgc | ggcccacctg | ctgggcccgc | 1260 |
| cccaccaccc | gggcctgccg | cctgaggccc | acctgaagcc | ggaacaccac | tacgccttca | 1320 |
| accacccgtt | ctccatcaac | aacctcatgt | cctcggagca | gcagcaccac | cacagccacc | 1380 |
| accaccacca | accccacaaa | atggacctca | aggcctacga | acaggtgatg | cactaccccg | 1440 |
| gctacggttc | ccccatgcct | ggcagcttgg | ccatgggccc | ggtcacgaac | aaaacgggcc | 1500 |
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| ttatgaactc | ctcttaagaa | gacgacggct | tcaggcccgg | ctaactctgg | caccccggat | 1620 |
| cgaggacaag | tgagagagca | agtgggggtc | gagactttgg | ggagacggtg | ttgcagagac | 1680 |
| | | | | | cagtcttctt | 1740 |
| cacccgctgc | agccgttccg | tcccaaacag | agggccacac | agatacccca | cgttctatat | 1800 |
| | | | | | | |

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aaqqaqqaaa acgggaaaga atataaagtt aaaaaaaagc ctccggtttc cactactgtg
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                                                                    1920
ccattgctgt tgttgcaggg aagtcttact taaaaaaaaa aaaaaatttt gtgagtgact
                                                                    1980
cggtgtaaaa ccatgtagtt ttaacagaac cagagggttg tactattgtt taaaaacagg
                                                                    2040
aaaaaaaata atqtaaqqqt ctqttgtaaa tgaccaaqaa aaagaaaaaa aaagcattcc
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                                                                    2242
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<210> 63

<211> 1047

<212> DNA

<213> NM_033423.2 \mid Homo sapiens granzyme H (cathepsin G-like 2, protein h-CCPX) (GZMH), mRNA

| <400> 63 | | | | | | |
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| ctcctgttgg | cctttcttct | gacccctggg | gctgggacag | aggagatcat | cgggggccat | 120 |
| gaggccaagc | cccactcccg | cccctacatg | gcctttgttc | agtttctgca | agagaagagt | 180 |
| cggaagaggt | gtggcggcat | cctagtgaga | aaggactttg | tgctgacagc | tgctcactgc | 240 |
| cagggaagct | ccataaatgt | caccttgggg | gcccacaata | tcaaggaaca | ggagcggacc | 300 |
| cagcagttta | tccctgtgaa | aagacccatc | ccccatccag | cctataatcc | taagaacttc | 360 |
| tccaacgaca | tcatgctact | gcagctggag | agaaaggcca | agtggaccac | agctgtgcgg | 420 |
| cctctcaggc | tacctagcag | caaggcccag | gtgaagccag | ggcagctgtg | cagtgtggct | 480 |
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| gtgcagaagg | actgccagtg | tgaacgtctc | ttccatggca | attacagcag | agccactgag | 600 |
| atttgtgtgg | gggatccaaa | gaagacacag | accggtttca | agggggactc | cggggggccc | 660 |
| ctcgtgtgta | aggacgtagc | ccaaggtatt | ctctcctatg | gaaacaaaaa | agggacacct | 720 |
| ccaggagtct | acatcaaggt | ctcacacttc | ctgccctgga | taaagagaac | aatgaagcgc | 780 |
| ctctaacagc | aggcatgaga | ctaaccttcc | tctgggcctg | accatctctg | ggacagaggc | 840 |
| aagaatcccc | aaggggtggg | cagtcagggt | tgcaggactg | taataaatgg | atctctggtg | 900 |
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<210> 64

<211> 5243

<212> DNA

<213> NM_001165.3| Homo sapiens baculoviral IAP repeat-containing 3 (BIRC3), transcript variant 1, mRNA

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| tccatgttct | agccaagtat | actattagaa | taaaaaaact | taacattgag | ttgcttcaac | 1740 |
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| agcatgaaac | tgagtccaaa | agaccaaatg | aacaaacaca | ttaatctctg | attatttatt | 1800 |
| ttaaatagaa | tatttaattg | tgtaagatct | aatagtatca | ttatacttaa | gcaatcatat | 1860 |
| tcctgatgat | ctatgggaaa | taactattat | ttaattaata | ttgaaaccag | gttttaagat | 1920 |
| gtgttagcca | gtcctgttac | tagtaaatct | ctttatttgg | agagaaattt | tagattgttt | 1980 |
| tgttctcctt | attagaagga | ttgtagaaag | aaaaaaatga | ctaattggag | aaaaattggg | 2040 |
| gatatatcat | atttcactga | attcaaaatg | tcttcagttg | taaatcttac | cattatttta | 2100 |
| cgtacctcta | agaaataaaa | gtgcttctaa | ttaaaatatg | atgtcattaa | ttatgaaata | 2160 |
| cttcttgata | acagaagttt | taaaatagcc | atcttagaat | cagtgaaata | tggtaatgta | 2220 |
| ttattttcct | cctttgagtt | aggtcttgtg | ctttttttc | ctggccacta | aatttcacaa | 2280 |
| tttccaaaaa | gcaaaataaa | catattctga | atatttttgc | tgtgaaacac | ttgacagcag | 2340 |
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| tgccactgaa | acattctagt | agcctggaga | agttgaccta | cctgtggaga | tgcctgccat | 2460 |
| taaatggcat | cctgatggct | taatacacat | cactcttctg | tgaagggttt | taattttcaa | 2520 |
| cacagcttac | tctgtagcat | catgtttaca | ttgtatgtat | aaagattata | caaaggtgca | 2580 |
| attgtgtatt | tcttccttaa | aatgtatcag | tataggattt | agaatctcca | tgttgaaact | 2640 |
| ctaaatgcat | agaaataaaa | ataataaaaa | atttttcatt | ttggcttttc | agcctagtat | 2700 |
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| ggataactgg | aaaagaggag | acagtcctac | tgaaaagcat | aaaaagttgt | atcctagctg | 3060 |
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| tttccgtggc | tcttattcaa | actctccatc | aaatcctgta | aactccagag | caaatcaaga | 3240 |
| tttttctgcc | ttgatgagaa | gttcctacca | ctgtgcaatg | aataacgaaa | atgccagatt | 3300 |
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| cttttactac | ataggacctg | gagacagagt | ggcttgcttt | gcctgtggtg | gaaaattgag | 3420 |
| caattgggaa | ccgaaggata | atgctatgtc | agaacacctg | agacattttc | ccaaatgccc | 3480 |
| atttatagaa | aatcagcttc | aagacacttc | aagatacaca | gtttctaatc | tgagcatgca | 3540 |
| gacacatgca | gcccgcttta | aaacattctt | taactggccc | tctagtgttc | tagttaatcc | 3600 |
| tgagcagctt | gcaagtgcgg | gtttttatta | tgtgggtaac | agtgatgatg | tcaaatgctt | 3660 |
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| ttgctgtgat | ggtggactca | ggtgttggga | atctggagat | gatccatggg | ttcaacatgc | 3720 |
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| agttcaagcc | agttaccctc | atctacttga | acagctgcta | tccacatcag | acagcccagg | 3840 |
| agatgaaaat | gcagagtcat | caattatcca | ttttgaacct | ggagaagacc | attcagaaga | 3900 |
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| caatgatctt | gtgttagact | tactcaatgc | agaagatgaa | ataagggaag | aggagaga | 4080 |
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| tattaatgaa | caagaacatg | atgttattaa | acagaagaca | cagacgtctt | tacaagcaag | 4260 |
| agaactgatt | gatacgattt | tagtaaaagg | aaatattgca | gccactgtat | tcagaaactc | 4320 |
| tctgcaagaa | gctgaagctg | tgttatatga | gcatttattt | gtgcaacagg | acataaaata | 4380 |
| tattcccaca | gaagatgttt | cagatctacc | agtggaagaa | caattgcgga | gactacaaga | 4440 |
| agaaagaaca | tgtaaagtgt | gtatggacaa | agaagtgtcc | atagtgttta | ttccttgtgg | 4500 |
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| tacaatcaag | ggtacagttc | gtacatttct | ttcatgaaga | agaaccaaaa | catcgtctaa | 4620 |
| actttagaat | taatttatta | aatgtattat | aactttaact | tttatcctaa | tttggtttcc | 4680 |
| ttaaaatttt | tatttattta | caactcaaaa | aacattgttt | tgtgtaacat | atttatatat | 4740 |
| gtatctaaac | catatgaaca | tatattttt | agaaactaag | agaatgatag | gcttttgttc | 4800 |
| ttatgaacga | aaaagaggta | gcactacaaa | cacaatattc | aatcaaaatt | tcagcattat | 4860 |
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| cccaggctga | ggcaagagaa | ttacttgagc | ccaggagttt | gaatccatcc | tgggcagcat | 5040 |
| actgagaccc | tgcctttaaa | aacaaacaga | acaaaaacaa | aacaccaggg | acacatttct | 5100 |
| ctgtctttt | tgatcagtgt | cctatacatc | gaaggtgtgc | atatatgttg | aatgacattt | 5160 |
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<210> 65

<400> 65

<211> 3850

<212> DNA

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|------------|-------------|------------|------------|------------|------------|------|
| aggccggtga | gggacctgcc | cagacctgga | gggtctcgct | ctgtcacaca | ggctggagtg | 120 |
| cagtggtgtg | atcttggctc | atcgtaacct | ccacctcccg | ggttcaagtg | attctcatgc | 180 |
| ctcagcctcc | cgagtagctg | ggattacagg | tggtgacttc | caagagtgac | tccgtcggag | 240 |
| gaaaatgact | ccccagtcgc | tgctgcagac | gacactgttc | ctgctgagtc | tgctcttcct | 300 |
| ggtccaaggt | gcccacggca | ggggccacag | ggaagacttt | cgcttctgca | gccagcggaa | 360 |
| ccagacacac | aggag cagcc | tccactacaa | acccacacca | gacctgcgca | tctccatcga | 420 |
| gaactccgaa | gaggccctca | cagtccatgc | ccctttccct | gcagcccacc | ctgcttcccg | 480 |
| atccttccct | gaccccaggg | gcctctacca | cttctgcctc | tactggaacc | gacatgctgg | 540 |
| gagattacat | cttctctatg | gcaagcgtga | cttcttgctg | agtgacaaag | cctctagcct | 600 |
| cctctgcttc | cagcaccagg | aggagagcct | ggctcagggc | ccccgctgt | tagccacttc | 660 |
| tgtcacctcc | tggtggagcc | ctcagaacat | cagcctgccc | agtgccgcca | gcttcacctt | 720 |
| ctccttccac | agtcctcccc | acacggccgc | tcacaatgcc | tcggtggaca | tgtgcgagct | 780 |
| caaaagggac | ctccagctgc | tcagccagtt | cctgaagcat | ccccagaagg | cctcaaggag | 840 |
| gccctcggct | gcccccgcca | gccagcagtt | gcagagcctg | gagtcgaaac | tgacctctgt | 900 |
| gagattcatg | ggggacatgg | tgtccttcga | ggaggaccgg | atcaacgcca | cggtgtggaa | 960 |
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| ccggagcggg | gaggctgaga | agagactcct | cctggtggac | ttcagcagcc | aagccctgtt | 1140 |
| ccaggacaag | aattccagcc | aagtcctggg | tgagaaggtc | ttggggattg | tggtacagaa | 1200 |
| caccaaagta | gccaacctca | cggagcccgt | ggtgctcact | ttccagcacc | agctacagcc | 1260 |
| gaagaatgtg | actctgcaat | gtgtgttctg | ggttgaagac | cccacattga | gcagcccggg | 1320 |
| gcattggagc | agtgctgggt | gtgagaccgt | caggagagaa | acccaaacat | cctgcttctg | 1380 |
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| gagcgccatg | ggctggggct | tccccatctt | tctggtgacg | ctggtggccc | tggtggatgt | 1860 |
| ggacaactat | ggccccatca | tcttggctgt | gcataggact | ccagagggcg | tcatctaccc | 1920 |
| ttccatgtgc | tggatccggg | actccctggt | cagctacatc | accaacctgg | gcctcttcag | 1980 |
| cctggtgttt | ctgttcaaca | tggccatgct | agccaccatg | gtggtgcaga | tcctgcggct | 2040 |

| gcgcccccac | acccaaaagt | ggtcacatgt | gctgacactg | ctgggcctca | gcctggtcct | 2100 |
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| gaagagaatc | aagaagaaaa | ataaaaatca | gctgttgtaa | tcacctagca | aactggaaaa | 3780 |
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| aaaaaaaaa | | | | | | 3850 |

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<211> 372

<212> DNA

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360

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| atcatctctc | gggagcaggc | ggatgagctt | cttggaggcg | tggagggtgc | ctacatcctt | 480 |
| agagaaagcc | agcggcaacc | aggatgctac | acgctggctc | tcaggtttgg | aaaccagacc | 540 |
| ttaaactaca | ggctcttcca | cgacgggaaa | cactttgtgg | gtgagaagag | gtttgagtcg | 600 |
| attcatgatc | tggtgacaga | tggcttgata | acactgtaca | tagaaacaaa | agctgccgag | 660 |
| tacatttcaa | aaatgacaac | taaccccatc | tatgaacaca | ttggatatgc | caccctactc | 720 |
| agagaaaaag | tatccagaag | gctgagcagg | tctaaaaatg | aaccaagaaa | aacaaacgtc | 780 |
| acacatgaag | aacacacagc | ggtggaaaag | atctcctccc | tggttcgaag | ggctgccctc | 840 |
| acacacaacg | acaaccactt | caattatgag | aagacacaca | actttaaggt | ccacacgttc | 900 |
| cgaggcccac | actggtgtga | atattgtgcc | aatttcatgt | gggggctcat | cgcccaaggg | 960 |
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| acctccacgt | gagaacaagg | gtgaaggtga | gggaagcccc | tcaggttggg | tcttttgctg | 1860 |
| tgcctcctat | gtatgtctgg | tttgctggaa | gagtgattaa | tacatcttta | atttattaaa | 1920 |
| aaacaatgta | gacctttaaa | cttcagtctt | attgggaata | aaagggaact | taattcatac | 1980 |

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<211> 2201

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| ttacaggaaa | actccagtga | ttttcaatca | aacattgcat | aggtaaccaa | ggagcagtga | 1440 |
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| acaatatagg | agctgtgtct | actattaaaa | gtgaaacatt | ttggcatgtt | tgttaattct | 1980 |
| agtttcattt | aataacctgt | aaggcacgta | agtttaagct | tttttttt | ttaagttaat | 2040 |
| gggaaaaatt | tgagacgcaa | taccaatact | taggattttg | gtcttggtgt | ttgtatgaaa | 2100 |
| ttctgaggcc | ttgatttaaa | tctttcattg | tattgtgatt | tccttttagg | tatattgcgc | 2160 |
| taagtgaaac | ttgtcaaata | aatcctcctt | ttaaaaactg | С | | 2201 |

<210> 75

<211> 1895

<212> DNA

<213> NM_004046.4| Homo sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, alpha subunit, isoform 1, cardiac muscle (ATP5A1), nuclear gene encoding mitochondrial protein, transcript variant 2, mRNA

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| agaagtaccg | cctgcggagt | aactgcaaag | atgctgtccg | tgcgcgttgc | tgcggccgtg | 120 |
| gtccgcgccc | ttcctcggcg | ggccggactg | gtctccagaa | atgctttggg | ttcatctttc | 180 |
| attgctgcaa | ggaacttcca | tgcctctaac | actcatcttc | aaaagactgg | gactgctgag | 240 |
| atgtcctcta | ttcttgaaga | gcgtattctt | ggagctgata | cctctgttga | tcttgaagaa | 300 |
| actgggcgtg | tcttaagtat | tggtgatggt | attgcccgcg | tacatgggct | gaggaatgtt | 360 |
| caagcagaag | aaatggtaga | gttttcttca | ggcttaaagg | gtatgtcctt | gaacttggaa | 420 |
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| attggtgacc | gacagactgg | gaaaacctca | attgctattg | acacaatcat | taaccagaaa | 780 |

| cgtttcaatg | atggatctga | tgaaaagaag | aagctgtact | gtatttatgt | tgctattggt | 840 |
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| agtaaattag | ttccatttgt | aaaagggtta | ctctcatact | ccttatgtac | agaaatcaca | 1860 |
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<211> 1290

<212> DNA

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| ggcccaggtc | aacataggca | tccacggcag | catctctgcc | gaagccaagg | cctgtgagga | 960 |
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<210> 78

<211> 4623

<212> DNA

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| atcgtgaagc | cgctggaagg | ttccgccaca | cttcaccact | ggcagcagtt | ggcccaacct | 1800 |
| caccttgggg | gcatcctgga | ccccggccc | ggtgtggtca | ccaagggctt | ccggacgctg | 1860 |

| gatgttgacc | tggacgaagt | gtactgcctt | aacgactttg | aagaagatga | cacaggtgac | 1920 |
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| cacatttctc | tcccacgcct | agctacctcc | actccagttc | agcacccaga | gacctcaggt | 1980 |
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| ggagagggga | gaacaatgaa | ttggctctat | tttctctatt | gggaattaca | ggaccatttt | 4080 |
| gattcttaga | atgtaaaaag | catatcgcta | agtaaatcat | cctggaggtc | ccaagtagct | 4140 |
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| gtatgataga | aggaaattgg | gaaaacttgt | atgctaggca | cttttgtcca | gagcctgctg | 4440 |
| tcccatggag | aaaaagtttt | aagcactgaa | aaaatttgat | taatgtattt | aaatgtatta | 4500 |
| tttgaagcat | cattcacttg | ttgattttta | caatcccatg | tcttaaaaag | gatgaatcca | 4560 |
| tgttattgta | ttgtaaataa | tttagattat | taaaatggat | tgtttaaaaa | aaaaaaaaa | 4620 |
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<210> 79

<211> 2657

<212> DNA

 $<\!213\!>$ NM_017895.6| Homo sapiens DEAD (Asp-Glu-Ala-Asp) box polypeptide 27 (DDX27), mRNA

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| gacgaggaag | aggaggggcc | cattgtgctg | ggcagacgac | aaaaagcttt | ggggaagaac | 240 |
| cgcagtgctg | atttcaaccc | tgatttcgtt | ttcactgaga | aggaggggac | gtacgatggc | 300 |
| agctgggccc | tggctgatgt | catgagccaa | ctcaagaaga | agagggcagc | cactacatta | 360 |
| gatgagaaga | ttgagaaagt | tcgaaagaaa | aggaaaacag | aggataaaga | agccaagtct | 420 |
| gggaagttgg | aaaaggagaa | agaagcaaag | gaaggctctg | aaccaaagga | gcaggaagac | 480 |
| cttcaagaga | atgatgagga | aggctcagaa | gatgaagcct | cggagactga | ctactcatca | 540 |
| gctgatgaga | acatcctcac | caaagcagat | acactcaaag | taaaggatcg | gaagaagaag | 600 |
| aagaagaaag | gacaggaagc | aggaggattt | tttgaagatg | catctcagta | cgatgaaaac | 660 |
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| accatgctct | tctcggccac | catgacagac | gaggtgaaag | atctggcttc | tgtctccttg | 1260 |
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| gcggaggaaa | aagagatgca | gcagtcagaa | gcccagatca | atacagcaaa | gcggctcctg | 1920 |
| gagaagggga | aggaggcagt | ggtccaagag | cccgagagga | gctggttcca | gaccaaagaa | 1980 |
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| cttgctgatt | agctttcata | tgactatatt | aaatggaagt | atttttggga | aaagagaaac | 2640 |
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<211> 3246

<212> DNA

<213> NM_018206.3| Homo sapiens vacuolar protein sorting 35 (yeast) (VPS35), mRNA

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<211> 3182

<212> DNA

<213> NM_017583.3| Homo sapiens tripartite motif-containing 44 (TRIM44), mRNA

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| ctgctactgc | cgccgccatg | ccgaggcgca | caggcagaag | ttcctcagtc | accatctggc | 480 |
| cgaatacgtc | cacggctccc | aggcctggac | cccgccagct | gacggagagg | gggcggggaa | 540 |
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| ccatgaatta | agatggatga | ctggaaaaag | gtgttggaga | aagagttaaa | gatgaggaag | 1920 |
| agatatttt | agtatatgaa | gttatccagg | acttgatatt | cataattcag | tgctgtggaa | 1980 |
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| tctcccacac | cctggatctc | actctcctct | tagtaacaga | gacactcctg | aggttggact | 2460 |
| tccttgcttt | tctctacttc | caaatcacaa | tttcttacaa | ccaagctttg | tgctcccgag | 2520 |
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| ctgttaaccc | agggcctaga | cttctagtgc | ctctgaggca | gaaccaaagg | agcctgcact | 2880 |
| ggggaaaatc | ccttttcctg | cctgcctgtc | tgcctgtgac | ctgtgtacgt | attacaggct | 2940 |
| ttaggaccag | ctgattgtta | tgcttgcagg | atggttttga | aacagaaaca | atacttgttt | 3000 |
| actgtaggaa | tcctatttat | attattttc | agtcctgtga | atgctgtgaa | aagatttatt | 3060 |
| cctttgaggc | caggaagctc | ccaggcatat | atgcttctag | gttaggattg | tcctgactca | 3120 |
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| aa | | | | | | 3182 |

<210> 82

<211> 4930

<212> DNA

 $<\!\!213\!\!>$ NM_020182.3| Homo sapiens transmembrane, prostate androgen induced RNA (TMEPAI), transcript variant 1, mRNA

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| ctgccagccc | attttccgga | cgccacccgc | gggcactgcc | gacgcccccg | gggctgccga | 180 |
| ggggaggccg | ggggggcgca | gcggagcgcg | gtcccgcgca | ctgagccccg | cggcgccccg | 240 |
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| ctgcatgcgg | ggccccagct | ccgggcgccg | gccggagccc | ccccggccg | ccccgagcc | 360 |
| cccgcgccc | cgcgccgcgc | cgccgcgccg | tccatgcacc | gcttgatggg | ggtcaacagc | 420 |
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ұссядстддс таладаттся ссудтдалау саудтусуд алстуссуся студассста
7460
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2400
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2280
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ST00
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1800
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T200
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1380
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T500
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1700
                 сссясстася дсудудсят судссястас ссудущест ссттссяцся ссядсядадс
1140
                 здтаастсуд усатсансяс саситиста высансяния институтительный возостительный возостительный возостительный высамостительный выпутывающий высамостительный выпутывающий высамостительный высамостите
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006
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840
                 ταςθεςεςεθε ετεθμέςεσες εφαισθεστίθη θε επίση ετεθρέσες το τροβουσιας το βεσθρομάτου το το διαθομό το το διαθομό το το διαθομό το διαθ
780
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120
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<211> 702

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<210> 84

<211> 2100

<212> DNA

<213> NM_015907.2| Homo sapiens leucine aminopeptidase 3 (LAP3), mRNA

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| cgcgcacgcc | gtctgcgccc | cgaaagcccc | gccccaaggc | gcgcccgccc | accgctctcc | 120 |
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| ccggctgcgg | ggcgagtagt | cgtccgacgt | ctggccgtga | gacgtttcgg | gagccggagt | 240 |
| ctctccaccg | cagacatgac | gaagggcctt | gttttaggaa | tctattccaa | agaaaaagaa | 300 |
| gatgatgtgc | cacagttcac | aagtgcagga | gagaattttg | ataaattgtt | agctggaaag | 360 |
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| tcttcactct | gtcttaaatt | ggacagttga | acttaaaagg | tttttgaata | aatggatgaa | 1800 |
| aatcttttaa | cggagacaaa | ggatggtatt | taaaaatgta | gaacacaatg | aaatttgtat | 1860 |
| gccttgattt | ttttttcatt | tcacacaaag | atttataaag | gtaaagttaa | tatcttactt | 1920 |
| gataaggatt | tttaagatac | tctataaatg | attaaaattt | ttagaacttc | ctaatcactt | 1980 |
| ttcagagtat | atgtttttca | ttgagaagca | aaattgtaac | tcagatttgt | gatgctagga | 2040 |

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<211> 1510

<212> DNA

<213> NM_018478.1 \mid Homo sapiens chromosome 20 open reading frame 35 (C20orf35), mRNA

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<211> 3105

<212> DNA

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MRNA
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| tttatgtcgt | cataccgtgg | ttacctgcat | actcttggtt | gttatcaact | tgttggtgat | 1680 |
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| cttcataccc | tccatgaagg | atatttttgg | agtcgtagga | gttacatctg | ctaacatgct | 1740 |
| tattttcatt | cttccttcat | ctctttattt | aaaaatcaca | accaggatgg | agataaagga | 1800 |
| actcaaagaa | tttgggctgc | ccttttcttg | ggcctggggg | tgttgttctc | cttggtcagc | 1860 |
| attcccttgg | tcatctatga | ctgggcctgc | tcatcgagta | atggtgaagg | ccactgaaac | 1920 |
| ccgccgagaa | aaagaaacat | ccctgttgtc | tgctcagtca | agtccccaca | catcagcaat | 1980 |
| ctctcaccac | ttcttttgca | agtttacaga | agcaaacaga | aatgtacagg | atacttaaaa | 2040 |
| tggaataact | ttttggttgc | aaaacagaga | catggttcta | taatgcttca | tgtccctcca | 2100 |
| agatttgaga | tcaatttagg | gattgtgaaa | ttttttttc | aaatttcata | caatcatatt | 2160 |
| tcccagtact | tttcacaatc | attttttacc | catctaactc | tatgttttgt | ggcttcccgg | 2220 |
| tctcttagaa | ctttgaaaac | atgatataca | ataatgttta | tttattatac | atccagattc | 2280 |
| tgaaataatt | ttcctactga | tgttcagctc | acactatctg | taccttttta | gaagagaaaa | 2340 |
| gaatcttgaa | ttgtatatat | ttattttgct | ttacagaaaa | aaatggtttc | gtaaataatt | 2400 |
| tgcctatttt | gggtaacata | gcacatggag | ataatcatct | gaaagttata | gggcactgcc | 2460 |
| actgctgaat | cagagcatgc | ccaatatttg | aggtggctct | gatttcctgg | cagctgaact | 2520 |
| cgggtagtcc | agtggcctag | ctggtaccac | atctattccc | atccagagac | attctctggc | 2580 |
| aagtgttctc | agctgaaaag | tggttgggga | tgattcttac | cttggtaatt | aaatgaagct | 2640 |
| acacatttgg | gtaatctagc | aaatgaagta | ttttttccct | cttggcaact | tgtgtcagag | 2700 |
| ttactctggt | ctgagtcaac | tttcgctggg | gaaaacctat | ggaacctact | gcaaaaagat | 2760 |
| tġtccaaaat | gcctaaġaaa | atactcctct | gatgcattta | gccttcaacc | ctacctgtct | 2820 |
| tgctgaaggg | agaaaaatgt | tttagtacat | tataggccca | gcagctttta | ttcatgtcca | 2880 |
| ccagctagtt | gcacagagaa | tcatgtgtac | ctaactaagg | atgatctagg | ataagtaact | 2940 |
| cctgttttat | attgagtatt | ttagggaagt | ctttaaaaga | cttgttttat | atctataaat | 3000 |
| ctaggttatt | acaaatacaa | gaattttgta | ccttaaataa | gcctcatttc | tatttcttct | 3060 |
| tcattaattc | tccatctagt | cttgtgaaaa | aaaaaaaaa | aaaaa | | 3105 |

<210> 87

<211> 2711

<212> DNA

<213> NM_016028.4| Homo sapiens suppressor of variegation 4-20 homolog 1 (Drosophila) (SUV420H1), transcript variant 2, mRNA

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| gcagcggccg | ggcagcgggg | cctgggacgc | gccccgagga | ggagcggggc | ggcgcaggcg | 180 |
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| gagagaacat | tgaaagtatt | ctctaagcta | tttgaagaga | gtgactaaat | gcacctgggt | 240 |
| caggctgtct | gtgggtatga | agtggttggg | agaatccaag | aacatggtgg | tgaatggcag | 300 |
| gagaaatgga | ggcaagttgt | ctaatgacca | tcagcagaat | caatcaaaat | tacagcacac | 360 |
| ggggaaggac | accctgaagg | ctggcaaaaa | tgcagtcgag | aggaggtcga | acagatgtaa | 420 |
| tggtaactcg | ggatttgaag | gacagagtcg | ctatgtacca | tcctctggaa | tgtccgccaa | 480 |
| ggaactctgt | gaaaatgatg | acctagcaac | cagtttggtt | cttgatccct | atttaggttt | 540 |
| tcaaacacac | aaaatgaata | ctagcgcctt | tccttcgagg | agctcaaggc | atttttcaaa | 600 |
| atctgacagt | ttttctcaca | acaaccctgt | gagatttagg | cctattaaag | gaaggcagga | 660 |
| agaactaaag | gaagtaattg | aacgttttaa | gaaagatgaa | cacttggaga | aagccttcaa | 720 |
| atgtttgact | tcaggcgaat | gggcacggca | ctattttctc | aacaagaata | aaatgcagga | 780 |
| gaaattattc | aaagaacatg | tatttattta | tttgcgaatg | tttgcaactg | acagtggatt | 840 |
| tgaaatattg | ccatgtaata | gatactcatc | agaacaaaat | ggagccaaaa | tagttgcaac | 900 |
| aaaagagtgg | aaacgaaatg | acaaaataga | attactggtg | ggttgtattg | ccgaactttc | 960 |
| agaaattgag | gagaacatgc | tacttagaca | tggagaaaac | gacttcagtg | tcatgtactc | 1020 |
| cacaaggaaa | aactgtgctc | aactctggct | gggtcctgct | gcgtttataa | accatgattg | 1080 |
| cagacctaat | tgtaagtttg | tgtcaactgg | tcgagataca | gcatgtgtga | aggctctaag | 1140 |
| agacattgaa | cctggagaag | aaatttcttg | ttattatgga | gatgggttct | ttggagaaaa | 1200 |
| taatgagttc | tgcgagtgtt | acacttgcga | aagacggggc | actggtgctt | ttaaatccag | 1260 |
| agtgggactg | cctgcgcctg | ctcctgttat | caatagcaaa | tatggactca | gagaaacaga | 1320 |
| taaacgttta | aataggctta | aaaagttagg | tgacagcagc | aaaaattcag | acagtcaatc | 1380 |
| tgtcagctct | aacactgatg | cagataccac | tcaggaaaaa | aacaatgcaa | gtaagtaagg | 1440 |
| gagatttgat | aagcatatct | tttaaaagta | ttttcacaca | atttgcttta | taaagtgtgc | 1500 |
| ttcagtagtt | ttaaactttt | aaatactcag | agagactggg | acttgtgagc | tttggctgca | 1560 |
| cttcaaggct | ctagacgtga | tttgagtaga | ggcacagtct | gtatcccatc | tctaacttca | 1620 |
| gtaccgtcct | ctagactatt | tttcttgaat | accttggtaa | ctggatatga | gttcttcatc | 1680 |
| atatgttcca | aggtcatcat | atgttttaaa | cattttcaag | gtgttagaga | ctgtgatgat | 1740 |
| gtcgctaagt | cctgcaagaa | gacaaaagga | ctgagtagaa | ttaaattaga | ctctatacat | 1800 |
| tccagtgcct | agccagtttg | ttagaaaaga | tgatggactt | ggggaattca | tagcttctgg | 1860 |
| ccttaaggct | tccacctttt | cattgcttgc | tgaccttttt | caaaacgaac | tgactcagtt | 1920 |
| cagcagacca | ccagtaccag | actcagaatt | gtgatagagg | agcattttga | acagtgccgt | 1980 |
| attgtgacat | gctgtattgg | ctactccaga | aagtaggagt | aaagatggaa | aggagaaaga | 2040 |
| agcaacctct | gagattccag | tggtgtgtgg | gggcaagatc | tgatggaaac | tgaaaaagag | 2100 |
| aacgaagact | aaacaaagag | aaaggaaaga | gaagaaaccc | taaatgggca | aaggaaagca | 2160 |
| | | | | | | |

| catcctgttt | gcggagcttt | gaaatattgg | aaccatttct | aattgctcct | gtttttctgg | 2220 |
|------------|------------|------------|------------|------------|------------|------|
| gtaacaccag | ttttctgtag | ttgccactaa | agcagtagac | tcttgagtct | cacttgtctc | 2280 |
| tgagagagac | agaagttaga | aagttttgac | ttggcgattc | cgaaagtatg | cctttgttgg | 2340 |
| cacttaaatg | tccagtgaga | cttcttggca | ccttagagcc | ctctgagata | ctgattattt | 2400 |
| taggttcttc | tccctacttt | cagatgtttt | cagcccaaca | ctgggtgctc | tcttccacta | 2460 |
| cagagaatcc | tgaagaaaag | ggaaggtgtt | tcccatgatg | gtgaatgtca | ctgccatgaa | 2520 |
| ttcctgaatc | tacctgctgc | tgggagtcag | agtccaagca | taacccgtgt | agcataaaag | 2580 |
| cagcgctgta | gccctattcc | agtctttttc | gttaatgtcc | agagtgaaca | acaagagtta | 2640 |
| gtcaatcatt | aactgttgac | tgttgattct | cataataaat | gcagcataac | gacaaaaaaa | 2700 |
| aaaaaaaaaa | a | | | | | 2711 |

<210> 88

<211> 2977

<212> DNA

<213> NM_022105.2| Homo sapiens death associated transcription factor 1 (DATF1), transcript variant 1, mRNA

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| cctcccggaa | gcgcggaacc | tcagcttccg | tacttgcgca | gaactcccct | cgcggcgacc | 180 |
| acgcactacg | ggttggcgcc | agagtcaaaa | ggcgtcggcc | ctctggcaag | atggctgctg | 240 |
| | | aatctggaac | | | | 300 |
| gcgaaggagc | ttactccacg | ggaacagcct | ctagataatc | tgagttgttg | aaaatacgaa | 360 |
| | | ggctgacaac | | | | 420 |
| | | cagggttttt | | | | 480 |
| | | caggtatgga | | | | 540 |
| | | ccagcaaaga | | | | 600 |
| | | gcgcagggga | | | | 660 |
| | | tgtccctgcg | | | | 720 |
| gcgcgtggag | cagttcctga | ccattgcgcg | gcgccgcggc | aggaggagca | tgcctgtctc | 780 |
| | | ccacgtcctg | | | | 840 |
| | | ctgagaccag | | | | 900 |
| | | ctgaaaaggt | | | | 960 |
| cgatagtgac | agcgatggcc | tgaccttgaa | agagcttcag | aatcgccttc | gcaggaagcg | 1020 |

| ggaacaggag | cccactgaga | ggcccctgaa | agggatccag | agtcgcctgc | ggaagaagcg | 1080 |
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| ccgggaggag | ggtcccgccg | agactgtggg | ctccgaggcc | agtgacactg | tggagggcgt | 1140 |
| cctgcccagt | aagcaggagc | ccgagaacga | tcagggggtt | gtgtcccagg | ctgggaaaga | 1200 |
| tgacagagag | agtaagttgg | agggaaaggc | ggctcaggac | atcaaagatg | aggagcctgg | 1260 |
| agacttgggc | cgaccgaagc | ctgaatgtga | gggttacgac | cccaacgccc | tgtattgcat | 1320 |
| ttgccgccag | cctcacaaca | acaggtttat | gatttgctgt | gaccgctgtg | aagaatggtt | 1380 |
| tcatggcgat | tgtgtgggca | tttctgaggc | tcgagggagg | cttttggaaa | ggaatgggga | 1440 |
| agactatatc | tgcccaaact | gcaccattct | gcaagtgcag | gatgagactc | attcagaaac | 1500 |
| ggcagatcag | caggaagcta | aatggagacc | tggagatgct | gatggcaccg | attgtacaag | 1560 |
| tataggaaca | atagagcaga | agtctagcga | agaccaaggg | ataaagggta | gaattgagaa | 1620 |
| agctgcaaat | ccaagtggca | agaagaaact | caagatcttc | cagcctgtga | tagaggcgcc | 1680 |
| tggtgcctca | aaatgtattg | gccccgggtg | ctgtcacgtg | gcgcagcccg | actcggtgta | 1740 |
| ctgcagtaat | gactgtatcc | tcaaacacgc | cgcagcgaca | atgaagtttc | taagctcagg | 1800 |
| taaagaacag | aagccaaagc | ctaaagaaaa | gatgaagatg | aagccagaga | agcccagtct | 1860 |
| tccgaaatgc | ggtgctcagg | caggtattaa | aatctcttct | gtgcacaaga | gaccagctcc | 1920 |
| agaaaaaaaa | gagaccacag | tgaagaaggc | agtggtggtc | cctgcgcgga | gtgaagcact | 1980 |
| cgggaaggaa | gcagcttgtg | agagcagcac | gccgtcgtgg | gcgagcgatc | acaattacaa | 2040 |
| tgcagtaaag | ccagaaaaga | ctgctgctcc | ctcgccgtca | ctgttgtata | aatgtatgta | 2100 |
| tcacctaggg | gttggcctcc | tggacccctc | ccgttctttc | tggatagcca | tcccctgggc | 2160 |
| ctgtccagga | ctgggagttg | cagctttgtg | ttaagctgat | cacagacacc | ggctgcacca | 2220 |
| tcagcgggaa | gcagagccca | tgtccaggat | gcctcctgct | gccctgtgtc | catccctagt | 2280 |
| ctgtcaggac | ttcctgtcac | tgttttccaa | agctgtaaac | ctcactggtg | aacgttcacc | 2340 |
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| tcttcatccc | agtctgattg | catagccaca | ctgcccggca | cgccacatcc | acccctgtct | 2460 |
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| gccagcacat | gaaagcatca | cttcttttt | atgttgtggg | aatctttgca | agttagtgtt | 2580 |
| gcatctgatt | ttcaggtgta | catttatttt | tgactgggca | gataggggat | tttttttt | 2640 |
| ccatgtccga | ttcacacgct | acacacccac | atgaacacat | tcgaacttcg | aaggccacac | 2700 |
| actcctgctt | cataggcccc | acggtaagtg | agttcacacc | tagaacactg | tcctgaccgc | 2760 |
| aggacgcgtg | ccttggactt | ggtattctac | atgtgactgg | ctttcttgcc | ctcgtctctt | 2820 |
| gaatgtttag | actcttaaga | tcatatcctg | ccccaaattt | caaattaatg | aaatgaagat | 2880 |
| atttcaaaca | gatctttgaa | acctcagatt | ctgtggtgca | attttaatgt | tttcttgttt | 2940 |
| ctcagttttc | tgctataaaa | ctattttcaa | ttcagtc | | | 2977 |
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<212> DNA

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agtatcttcg ttcggtgaaa gaaacacctc atcgtccatc agacgggctt tcaaataccg 180

| agtcttcgga | tgggttgaat | aagctacttg | ctcatctgct | tatgctttct | aagaggtgtc | 240 |
|------------|------------|------------|------------|------------|------------|------|
| ccttcaaaga | tgtgagagag | aaaagtgagt | ttattctgaa | gagcatccag | gaacttggca | 300 |
| ttagaattcc | tcgaccacta | ggacagggac | caagcagatt | catcccagaa | aaggagatcc | 360 |
| tccaagtggg | gagtgaagac | gcacagatgc | atgctttatt | tgcagattct | tttgctgctt | 420 |
| tgggccgttt | ggataacatt | acgttagtga | tggttttcca | cccacaatat | ttagaaagtt | 480 |
| tcttaaaaac | tcagcactat | ctactgcaaa | tggatgggcc | gttaccccta | cattatcgtc | 540 |
| actacattgg | aataatggct | gcggcaagac | atcagtgctc | ctacttagtg | aacctgcatg | 600 |
| taaatgattt | ccttcatgtt | ggtggggacc | ccaagtggct | caatggttta | gagaatgctc | 660 |
| ctcaaaaact | acagaattta | ggagaactta | acaaagtgtt | agcccataga | ccttggctta | 720 |
| ttaccaaaga | acacattgag | ggacttttaa | aagctgaaga | gcacagctgg | tcccttgcgg | 780 |
| aattggtaca | tgcagtagtt | ttactcacac | actatcattc | tcttgcctca | ttcacattcg | 840 |
| gctgtggaat | cagtccagaa | attcattgtg | atggtggcca | cacattcaga | cctccttctg | 900 |
| ttagcaacta | ctgcatctgt | gacattacaa | atggcaatca | cagtgtggat | gagatgccgg | 960 |
| tcaactcagc | agaaaatgtt | tctgtaagtg | attctttctt | tgaggttgaa | gccctcatgg | 1020 |
| aaaagatgag | gcagttacag | gaatgtcgag | atgaagaaga | ggcaagtcag | gaagagatgg | 1080 |
| cttcacgttt | tgaaatagaa | aaaagagaga | gtatgtttgt | cttctcttca | gatgatgaag | 1140 |
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| atttctctag | acatgggatg | catgttccaa | catttcgtgt | ccaggactat | tgctgggaag | 1260 |
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| aatttcacat | tgcttacaat | cttacttata | atacaatggc | aatgcacaaa | gatgttgata | 1380 |
| cctcaatgct | tagacgggca | atttggaact | atattcactg | catgtttgga | ataagatatg | 1440 |
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| ttaaagatga | ttctggaatg | atcagcagat | atagtctaca | agggggaagg | tactaagccc | 1740 |
| caggaccaat | ggtagacaaa | ataattcaga | aatccattgt | gccatgattc | ctttagtttc | 1800 |
| tgctatttt | ctgtggaaaa | ccactgctgg | cacaagcagt | gactgtttgg | cagcttcaag | 1860 |
| tttagagctg | tgaagacagg | ctgccattca | cagtattttg | ctttttgaca | gtacaagatg | 1920 |
| ctgtgtaact | gttttaatac | agcaaatagt | aactctccaa | atcctgttgc | ttttatgtta | 1980 |
| aataagataa | caagaattgg | agcatgcaaa | gaatgggact | tggataatga | cttaagcttt | 2040 |
| atatgtaaag | aattttagaa | gatcttggtg | ctgctattcc | tgctggagga | atgaatagat | 2100 |
| ggctgtttca | gttaagctat | tagtaataaa | agtgaacatt | gctactatct | gagcctacat | 2160 |
| acataacttg | tgtgatttca | aattaaactt | gcattatgtg | ttaattttct | tgcatctaaa | 2220 |

| aaagcataga attcctactc | acacagctca | gcaacaacca | ttttgatggt | aacagttaat | 2280 |
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| ttctttcatt agttttttaa | attcagggtt | ctggatatta | aattaaaatg | gcattcttaa | 2340 |
| agattttctt caaaaagcaa | tcctaaatga | aagtgtgtaa | attataagaa | gctggcgatc | 2400 |
| ttttgatatg ctgtttcaca | ggatcctgac | actggagggc | agctgtcttg | tgcattactt | 2460 |
| gtgttcccag caccaaagtt | gtgggacatg | ttgctgtaga | ctgctgcgca | gtcctgggtg | 2520 |
| cattcagtct ctctgcctct | gcctgcctcc | tggtccccac | tttaaaggct | gtgcagctcc | 2580 |
| ttaaataata aagctggaaa | atatttttag | tcgggttatc | aaatttgatt | tacaaaaacg | 2640 |
| ctaactttgt ttgaaatgca | aacaggtttg | aaaatatgta | ttaagtactt | tgtattctgg | 2700 |
| aagcgtgaat tgcttttgaa | gtctgtcagt | attactggta | tttttaaata | aagaagaatt | 2760 |
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<210> 91

<211> 3802

<212> DNA

<213> NM_017763.1| Homo sapiens hypothetical protein FLJ20315 (FLJ20315), mRNA

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| aagcataaat | gttcttttcc | tccatttgtc | tggatctgag | aacctgcatt | tggtattagc | 120 |
| tagtggaagc | agtatgtatg | gttgaagtgc | attgctgcag | ctggtagcat | gagtggtggc | 180 |
| caccagctgc | agctggctgc | cctctggccc | tggctgctga | tggctaccct | gcaggcaggc | 240 |
| tttggacgca | caggactggt | actggcagca | gcggtggagt | ctgaaagatc | agcagaacag | 300 |
| aaagctgtta | tcagagtgat | ccccttgaaa | atggacccca | caggaaaact | gaatctcact | 360 |
| ttggaaggtg | tgtttgctgg | tgttgctgaa | ataactccag | cagaaggaaa | attaatgcag | 420 |
| tcccacccac | tgtacctgtg | caatgccagt | gatgacgaca | atctggagcc | tggattcatc | 480 |
| agcatcgtca | agctggagag | tcctcgacgg | gcccccgcc | cctgcctgtc | actggctagc | 540 |
| aaggctcgga | tggcgggtga | gcgaggagcc | agtgctgtcc | tctttgacat | cactgaggat | 600 |
| cgagctgctg | ctgagcagct | gcagcagccg | ctggggctga | cctggccagt | ggtgttgatc | 660 |
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| aggattgagc | tgaaggagcc | cccggcctgg | ccagattatg | atgtgtggat | cctaatgaca | 780 |
| gtggtgggca | ccatctttgt | gatcatcctg | gcttcggtgc | tgcgcatccg | gtgccgcccc | 840 |
| cgccacagca | ggccggatcc | gcttcagcag | agaacagcct | gggccatcag | ccagctggcc | 900 |
| accaggaggt | accaggccag | ctgcaggcag | gcccggggtg | agtggccaga | ctcagggagc | 960 |
| agctgcagct | cagcccctgt | gtgtgccatc | tgtctggagg | agttctctga | ggggcaggag | 1020 |

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| tccctgggac | cctctcgatc | ttaccaagaa | ccaggtcgaa | gactccacct | cattcgccag | 1200 |
| catcccggcc | atgcccacta | ccacctccct | gctgcctacc | tgttgggccc | ttcccggagt | 1260 |
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| cctcggcatc | accgcttccc | cagagctgca | catccccggg | ctccaggaga | gcagcagcgc | 1380 |
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| tcacagcacc | ctgctgcttg | cccagtgccc | ctacgccggg | ccaggccccc | tgacagcagt | 1500 |
| ggatctggag | aaagctattg | cacagaacgc | agtgggtacc | tggcagatgg | gccagccagt | 1560 |
| gactccagct | cagggccctg | tcatggctct | tccagtgact | ctgtggtcaa | ctgcacggac | 1620 |
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| tttgaccccc | tagtgtactg | cagccctaaa | ggggatcccc | agcgagtgga | catgcagcct | 1740 |
| agtgtgacct | ctcggcctcg | ttccttggac | tcggtggtgc | ccacagggga | aacccaggtt | 1800 |
| tccagccatg | tccactacca | ccgccaccgg | caccaccact | acaaaaagcg | gttccagtgg | 1860 |
| catggcagga | agcctggccc | agaaaccgga | gtcccccagt | ccaggcctcc | tattcctcgg | 1920 |
| acacagcccc | agccagagcc | accttctcct | gatcagcaag | tcaccggatc | caactcagca | 1980 |
| gccccttcgg | ggcggctctc | taacccacag | tgccccaggg | ccctccctga | gccagcccct | 2040 |
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<211> 1236

<212> DNA

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| cgacccctgg | cacctggcgc | ccagcgcgcc | cgtggccgct | gccgcctccg | ccccaggttt | 180 |
| tgcgtgtgaa | gctgtgtgga | aatgtgaaat | actaccagtc | acaccattat | agtaccgtgg | 240 |
| tgccacctga | tgaaataaca | gttatttata | gacatggcct | tcccttggta | acacttacct | 300 |
| tgccatctag | aaaagaacgt | tgtcaattcg | tagtcaaacc | aatgttgtca | acagttggtt | 360 |
| cattccttca | ggacctacaa | aatgaagata | agggtatcaa | aactgcagcc | atcttcacag | 420 |
| cagatggcaa | catgatttca | gcttctacct | tgatggatat | tttgctaatg | aatgatttta | 480 |
| aacttgtcat | taataaaata | gcatatgatg | tgcagtgtcc | aaagagagaa | aaaccaagta | 540 |
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| tgcatttaga | agagtctcag | aaaaagagag | agcaccattt | actggagaaa | attgaccacc | 660 |
| tgaaggaaca | gctgcagccc | cttgaacagg | tgaaagctgg | aatagaagct | cattcggaag | 720 |
| ccaaaaccag | tggactcctg | tgggctggat | tggcactgct | gtccattcag | ggtggggcac | 780 |
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| tcacatttgc | aaattctatg | gtcttttttg | catactttat | agtcactcga | caggattata | 900 |
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| agaattaatc | ttacagtttt | aaatgtcgtc | agattttcca | ttatgtattg | attttgcaac | 1140 |
| ttaggatgtt | tttgagtccc | atggttcatt | ttgattgttt | aatctttgtt | attaaattct | 1200 |
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<210> 93

<211> 2096

<212> DNA

 $<\!\!213\!\!>$ NM_024792.1| Homo sapiens membrane protein expressed in epithelial-like lung adenocarcinoma (CT120), mRNA

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| agccc | ggatg | gagccgcacc | gactgcgtga | tgatcagcac | caggctggtt | tcctcggtgc | 180 |
| acgcc | gtgct | ggccaccggc | tcggggatcg | tcatcattcg | ctcctgcgac | gacgtgatca | 240 |
| ccggc | aggca | ctggcttgcc | cgggaatatg | tgtggtttct | gattccatac | atgatctatg | 300 |
| actcg | jtacgc | catgtacctc | tgtgaatggt | gccgaaccag | agaccagaac | cgtgcgccct | 360 |
| ccctc | actct | tcgaaacttc | ctaagtcgaa | accgcctcat | gatcacacat | catgcggtca | 420 |
| ttctc | tttgt | ccttgtgcca | gtcgcacaga | ggctccgggg | agaccttggg | gacttctttg | 480 |
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| ccttc | ctcgt | agctcctcag | atctactggt | tctgtctgct | gtgcaggaag | gcagtccggc | 780 |
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| tgtca | gacct | ccacggacag | caaagtggtt | ttaatgcaag | cccaaggatc | cttcttaagg | 1080 |
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| gacagtaccc | gtggcactgg | gcccgcagaa | gcaagggatg | acttggttct | tggaagtaat | 1260 |
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| tagataaccc | attggttctt | tgcctcatcc | tctcatccat | gggtcagagt | tgaattctta | 1380 |
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| caagcaggat | tttctggtgc | caacactcat | tcatcattcc | cgatcaacta | ggatgaattt | 1980 |
| aagactgtgc | taccatgtgt | tctcaagtgg | tagtttaaaa | agtggatttt | taaagtgcct | 2040 |
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<210> 94

<211> 4372

<212> DNA

<213> NM_014314.2| Homo sapiens DEAD (Asp-Glu-Ala-Asp) box polypeptide 58 (DDX58), mRNA

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| agggcccaat | ggaggctgcc | acactttttc | tcaagttcct | gttggagctc | caggaggaag | 360 |
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| aagacttctt | cagcaatgtc | cgagcagcag | gattcgatga | gattgagcaa | gatcttactc | 1920 |
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| aattcatcag | agatagtcaa | gaaaaaccaa | aacctgtacc | tgataaggaa | aataaaaaac | 2580 |
| tgctctgcag | aaagtgcaaa | gccttggcat | gttacacagc | tgacgtaaga | gtgatagagg | 2640 |
| aatgccatta | cactgtgctt | ggagatgctt | ttaaggaatg | ctttgtgagt | agaccacatc | 2700 |
| ccaagccaaa | gcagttttca | agttttgaaa | aaagagcaaa | gatattctgt | gcccgacaga | 2760 |
| | | | | | | |

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actgcagcca tgactgggga atccatgtga agtacaagac atttgagatt ccagttataa
                                                                    2820
                                                                    2880
aaattgaaag ttttgtggtg gaggatattg caactggagt tcagacactg tactcgaagt
ggaaggactt tcattttgag aagataccat ttgatccagc agaaatgtcc aaatgatatc
                                                                    2940
                                                                    3000
aggtcctcaa tcttcagcta cagggaatga gtaactttga gtggagaaga aacaaacata
gtgggtataa tcatggatcg cttgtacccc tgtgaaaata tatttttaa aaatatcttt
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tttgtaggcc aacagagctc atagtacttg ggaaaaatta aaaagcctca tttctagcct
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                                                                     3480
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tgtgaaaagt acatgacaca gtagttgctt gataatagtt actagtagta gtattcttac
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<210> 95

<211> 2163

<212> DNA

<213> NM_015515.3| Homo sapiens keratin 23 (histone deacetylase inducible) (KRT23), transcript variant 1, mRNA

| <400> 95 ggcagatgaa | atataagatt | catcaaccac | atttgacagc | ccatggcagg | tttcctgttt | 60 |
|------------------------|------------|------------|------------|------------|------------|------|
| tccatcgtcc | ctctgcaggt | cacagacaca | cagagcccag | ccgtggcagg | ctcagccggg | 120 |
| gtccggggct | gctaacaacg | gctacattcc | tccccaggg | ccaagggaaa | tcctgagcgc | 180 |
| aggccagggt | tgtttggttt | tgaggtgtgc | tgggatgaaa | ggcaccctgg | aagtggaagg | 240 |
| taaatgaaca | atggaaaaac | ttcacggcaa | gattagaaag | atacctgagc | ccaatacccg | 300 |
| cctgatgtcg | tgggccacac | ctccgggtta | ccaggggaag | ggaggaagca | aactgtcata | 360 |
| ttgatgtggc | tctaaacaac | aacagtgtgc | gaaggcccag | gggcactttg | ggattgacca | 420 |
| agaggaaaca | caagttgcac | aatgatacaa | tcttgttggt | acaattgtca | gagaagggaa | 480 |
| ctcccacagc | aaaggccata | aaaccatcca | gggcagtctg | gggcggctca | gttctgcggt | 540 |
| gccagggagt | ggagcagagc | tcagccccgt | cccaaacaca | gatgggacca | tgaactccgg | 600 |
| acacagcttc | agccagaccc | cctcggcctc | cttccatggc | gccggaggtg | gctggggccg | 660 |
| gcccaggagc | ttccccaggg | ctcccaccgt | ccatggcggt | gcgggggag | cccgcatctc | 720 |
| cctgtccttc | accacgcgga | gctgcccacc | ccctggaggg | tcttggggtt | ctggaagaag | 780 |
| cagcccccta | ctaggcggaa | atgggaaggc | caccatgcag | aatctcaacg | accgcctggc | 840 |
| ctcctacctg | gagaaggttc | gcgccctgga | ggaggccaac | atgaagctgg | aaagccgcat | 900 |
| cctgaaatgg | caccagcaga | gagatcctgg | cagtaagaaa | gattattccc | agtatgagga | 960 |
| aaacatcaca | cacctgcagg | agcagatagt | ggatggtaag | atgaccaatg | ctcagattat | 1020 |
| tcttctcatt | gacaatgcca | ggatggcagt | ggatgacttc | aacctcaagt | atgaaaatga | 1080 |
| acactccttt | aagaaagact | tggaaattga | agtcgagggc | ctccgaagga | ccttagacaa | 1140 |
| cctgaccatt | gtcacaacag | acctagaaca | ggaggtggaa | ggaatgagga | aagagctcat | 1200 |
| tctcatgaag | aagcaccatg | agcaggaaat | ggagaagcat | catgtgccaa | gtgacttcaa | 1260 |
| tgtcaatgtg | aaggtggata | caggtcccag | ggaagatctg | attaaggtcc | tggaggatat | 1320 |
| gagacaagaa | tatgagctta | taataaagaa | gaagcatcga | gacttggaca | cttggtataa | 1380 |
| agaacagtct | gcagccatgt | cccaggaggc | agccagtcca | gccactgtgc | agagcagaca | 1440 |
| aggtgacatc | cacgaactga | agcgcacatt | ccaggccctg | gagattgacc | tgcagacaca | 1500 |
| gtacagcacg | aaatctgctt | tggaaaacat | gttatccgag | acccagtctc | ggtactcctg | 1560 |
| caagctccag | gacatgcaag | agatcatctc | ccactatgag | gaggaactga | cgcagctacg | 1620 |
| ccatgaactg | gagcggcaga | acaatgaata | ccaagtgctg | ctgggcatca | aaacccacct | 1680 |
| ggagaaggaa | atcaccacgt | accgacggct | cctggaggga | gagagtgaag | ggacacggga | 1740 |
| agaatcaaag | tcgagcatga | aagtgtctgc | aactccaaag | atcaaggcca | taacccagga | 1800 |
| gaccatcaac | ggaagattag | ttctttgtca | agtgaatgaa | atccaaaagc | acgcatgaga | 1860 |
| ccaatgaaag | tttccgcctg | ttgtaaaatc | tattttcccc | caaggaaagt | ccttgcacag | 1920 |
| acaccagtga | gtgagttcta | aaagataccc | ttggaattat | cagactcaga | aacttttatt | 1980 |
| | | | | | | |

<210> 96

<211> 2881

<212> DNA

<213> NM_007210.2| Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 6 (GalNAc-T6) (GALNT6), mRNA

<400> 96 atgaggetee teegcagaeg ceacatgeee etgegeetgg ceatggtggg etgegeettt 60 gtgctcttcc tcttcctcct gcatagggat gtgagcagca gagaggaggc cacagagaag 120 ccgtggctga agtccctggt gagccggaag gatcacgtcc tggacctcat gctggaggcc 180 atgaacaacc ttagagattc aatgcccaag ctccaaatca gggctccaga agcccagcag 240 300 actctgttct ccataaacca gtcctgcctc cctgggttct ataccccagc tgaactgaag cccttctggg aacggccacc acaggacccc aatgcccctg gggcagatgg aaaagcattt 360 cagaagagca agtggacccc cctggagacc caggaaaagg aagaaggcta taagaagcac 420 tgtttcaatg cctttgccag cgaccggatc tccctgcaga ggtccctggg gccagacacc 480 540 cqaccacctq aqtqtqtgga ccagaagttc cggcgctgcc ccccactggc caccaccagc gtgatcattg tgttccacaa cgaagcctgg tccacactgc tgcgaacagt gtacagcgtc 600 660 ctacacacca cccctgccat cttgctcaag gagatcatac tggtggatga tgccagcaca 720 gaqqaqcacc taaaqqagaa gctggagcag tacgtgaagc agctgcaggt ggtgagggtg 780 gtgcggcagg aggagcggaa ggggttgatc accgcccggc tgctgggggc cagcgtggca 840 caggcggagg tgctcacgtt cctggatgcc cactgtgagt gcttccacgg ctggctggag cccctcctgg ctcgaatcgc tgaggacaag acagtggtgg tgagcccaga catcgtcacc 900 atcqacctta atacttttqa qttcqccaag cccgtccaga ggggcagagt ccatagccga 960 1020 ggcaactttg actggagcct gaccttcggc tgggaaacac ttcctccaca tgagaagcag 1080 aggcgcaagg atgaaacata ccccatcaaa tccccgacgt ttgctggtgg cctcttctcc 1140 atccccaagt cctactttga gcacatcggt acctatgata atcagatgga gatctgggga 1200 ggggagaacg tggaaatgtc cttccgggtg tggcagtgtg ggggccagct ggagatcatc 1260 ccctgctctg tcgtaggcca tgtgttccgg accaagagcc cccacacctt ccccaagggc 1320 actaqtqtca ttgctcgcaa tcaagtgcgc Ctggcagagg tctggatgga cagctacaag 1380 aagattttct ataggagaaa tctgcaggca gcaaagatgg cccaagagaa atccttcggt

| gacatttcgg | aacgactgca | gctgagggaa | caactgcact | gtcacaactt | ttcctggtac | 1440 |
|------------|------------|------------|------------|------------|------------|------|
| ctgcacaatg | tctacccaga | gatgtttgtt | cctgacctga | cgcccacctt | ctatggtgcc | 1500 |
| atcaagaacc | tcggcaccaa | ccaatgcctg | gatgtgggtg | agaacaaccg | cggggggaag | 1560 |
| cccctcatca | tgtactcctg | ccacggcctt | ggcggcaacc | agtactttga | gtacacaact | 1620 |
| cagagggacc | ttcgccacaa | catcgcaaag | cagctgtgtc | tacatgtcag | caagggtgct | 1680 |
| ctgggccttg | ggagctgtca | cttcactggc | aagaatagcc | aggtccccaa | ggacgaggaa | 1740 |
| tgggaattgg | cccaggatca | gctcatcagg | aactcaggat | ctggtacctg | cctgacatcc | 1800 |
| caggacaaaa | agccagccat | ggccccctgc | aatcccagtg | acccccatca | gttgtggctc | 1860 |
| tttgtctagg | acccagatca | tccccagaga | gagcccccac | aagctcctca | ggaaacagga | 1920 |
| ttgctgatgt | ctgggaacct | gatcaccagc | ttctctggag | gccgtaaaga | tggatttcta | 1980 |
| aacccactgg | gtggcaaggc | aggaccttcc | taatccttgc | aacaacattg | ggcccatttt | 2040 |
| ctttccttca | caccgatgga | agagaccatt | aggacatata | tttagcctag | cgttttcctg | 2100 |
| ttctagaaat | agaggctccc | aaagtaggga | aggcagctgg | gggagggttc | agggcagcaa | 2160 |
| tgctgagttc | aagaaaagta | cttcaggctg | ggcacagtgg | ctcatgcctg | aaatcctagc | 2220 |
| actttgggaa | gacaatgtgg | gagaatggct | tgagcccagg | agttcaagac | cggcctgagc | 2280 |
| aacatagtga | ggatcccatc | tctacgccca | ccctccccc | ggcaaaaaaa | aaagctgggt | 2340 |
| atggtggctt | atgcctgtag | tcgcagctac | tcagaaggct | gaggtgggag | gattgcttgt | 2400 |
| tccccggagg | ttgaagctac | agtgagcctt | gattgtgtca | ctgcactcca | gcctgggcaa | 2460 |
| caggtaagac | tctgtctcaa | aaaaaaaaca | aaaaagaaga | agaaaagtac | ttctacagcc | 2520 |
| atgtcctatt | ccttgatcat | ccaaagcacc | tgcagagtcc | agtgaaatga | tatattctgg | 2580 |
| ctgggcacag | tggctcacac | ctgtaatcct | agcactttgg | gaggccaagg | caggtggatc | 2640 |
| acctgaggtc | agaagtttga | aaccagcctg | gactacatgg | tgaaactcca | tctctactaa | 2700 |
| aagtacaaaa | attagctggg | catgatggca | cgcacctgca | gtcccagcta | cttgggaggc | 2760 |
| tgaggcagga | gaatcactcg | aacccaggag | gcagaggttg | cagtgagcca | agacagcacc | 2820 |
| attgcacccc | agcctgagca | acaagagcga | aactccatct | caggaaaaaa | aaaaaaaaa | 2880 |
| a | | | | | | 2881 |

<210> 97

<211> 1930

<212> DNA

<213> NM_020183.3| Homo sapiens aryl hydrocarbon receptor nuclear translocator-like 2 (ARNTL2), mRNA

<400> 97 gaccaagtgg ctcctgcgat ggcggcggaa gaggaggctg cggcgggagg taaagtgttg 60

| agagaggaga | accagtgcat | tgctcctgtg | gtttccagcc | gcgtgagtcc | agggacaaga | 120 |
|------------|------------|------------|------------|---------------------|------------|------|
| ccaacagcta | tggggtcttt | cagctcacac | atgacagagt | ttccacgaaa | acgcaaagga | 180 |
| agtgattcag | acccatccca | gtcaggaatc | atgacagaaa | aagt g gtgga | aaagctttct | 240 |
| cagaatcccc | ttacctatct | tctttcaaca | aggatagaaa | tatcagcctc | cagtggcagc | 300 |
| agagtggaag | atggtgaaca | ccaagttaaa | atgaaggcct | tcagagaagc | tcatagccaa | 360 |
| actgaaaagc | ggaggagaga | taaaatgaat | aacctgattg | aagaactgtc | tgcaatgatc | 420 |
| cctcagtgca | accccatggc | gcgtaaactg | gacaaactta | cagttttaag | aatggctgtt | 480 |
| caacacttga | gatctttaaa | aggcttgaca | aattcttatg | tgggaagtaa | ttatagacca | 540 |
| tcatttcttc | aggataatga | gctcagacat | ttaatcctta | agactgcaga | aggcttctta | 600 |
| tttgtggttg | gatgtgaaag | aggaaaaatt | ctcttcgttt | ctaagtcagt | ctccaaaata | 660 |
| cttaattatg | atcaggctag | tttgactgga | caaagcttat | ttgacttctt | acatccaaaa | 720 |
| gatgttgcca | aagtaaagga | acaactttct | tcttttgata | tttcaccaag | agaaaagcta | 780 |
| atagatgcca | aaactggttt | gcaagttcac | agtaatctcc | acgctggaag | gacacgtgtg | 840 |
| tattctggct | caagacgatc | tttttctgt | cggataaaga | gttgtaaaat | ctctgtcaaa | 900 |
| gaagagcatg | gatgcttacc | caactcaaag | aagaaagagc | acagaaaatt | ctatactatc | 960 |
| cattgcactg | gttacttgag | aagctggcct | ccaaatattg | ttggaatgga | agaagaaagg | 1020 |
| aacagtaaga | aagacaacag | taattttacc | tgccttgtgg | ccattggaag | attacagcca | 1080 |
| tatattgttc | cacagaacag | tggagagatt | aatgtgaaac | caactgaatt | tataacccgg | 1140 |
| tttgcagtga | atggaaaatt | tgtctatgta | gatcaaaggg | caacagcgat | tttaggatat | 1200 |
| ctgcctcagg | aacttttggg | aacttcttgt | tatgaatatt | ttcatcaaga | tgaccacaat | 1260 |
| aatttgactg | acaagcacaa | agcagttcta | cagagtaagg | agaaaatact | tacagattcc | 1320 |
| tacaaattca | gagcaaaaga | tggctctttt | gtaactttaa | aaagccaatg | gtttagtttc | 1380 |
| acaaatcctt | ggacaaaaga | actggaatat | attgtatctg | tcaa cacttt | agttttggga | 1440 |
| catagtgagc | ctggagaagc | atcattttta | ccttgtagct | ctcaatcatc | agaagaatcc | 1500 |
| tctagacagt | cctgtatgag | tgtacctgga | atgtctactg | gaacagtact | tggtgctggt | 1560 |
| agtattggaa | cagatattgc | aaatgaaatt | ctggatttac | agaggttaca | gtcttcttca | 1620 |
| taccttgatg | attcgagtcc | aacaggttta | atgaaagata | ctcatactgt | aaactgcagg | 1680 |
| agtatgtcaa | ataaggagtt | gtttccacca | agtccttctg | aaat gg ggga | gctagaggct | 1740 |
| accaggcaaa | accagagtac | tgttgctgtc | cacagccatg | agccactcct | cagtgatggt | 1800 |
| gcacagttgg | atttcgatgc | cctatgtgac | aatgatgaca | cagc catggc | tgcatttatg | 1860 |
| aattacttag | aagcagaggg | gggcctggga | gaccctgggg | actt cagtga | catccagtgg | 1920 |
| accctctagc | | | | | | 1930 |
| | | | | | | |

<210> 98

<211> 2128

<212> DNA

<213> NM_014576.2| Homo sapiens apobec-1 complementation factor (ACF), transcript variant 1, mRNA

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|------------------------|------------|------------|------------|------------|------------|------|
| accttttccg | ggtggggatc | tctgaaatta | ctcagataac | agtgctgtgc | caaaaacctg | 120 |
| tggattttct | ctacaaaaat | tattgagcaa | ccctaattaa | cctgattttt | tgctgataat | 180 |
| cactctcaat | ggaatcaaat | cacaaatccg | gggatggatt | gagcggcact | cagaaggaag | 240 |
| cagccctccg | cgcactggtc | cagcgcacag | gatatagctt | ggtccaggaa | aatggacaaa | 300 |
| gaaaatatgg | tggccctcca | cctggttggg | atgctgcacc | ccctgaaagg | ggctgtgaaa | 360 |
| tttttattgg | aaaacttccc | cgagaccttt | ttgaggatga | gcttatacca | ttatgtgaaa | 420 |
| aaatcggtaa | aatttatgaa | atgagaatga | tgatggattt | taatggcaac | aatagaggat | 480 |
| atgcatttgt | aacattttca | aataaagtgg | aagccaagaa | tgcaatcaag | caacttaata | 540 |
| attatgaaat | tagaaatggg | cgcctcttag | gggtttgtgc | cagtgtggac | aactgccgat | 600 |
| tatttgttgg | gggcatccca | aaaaccaaaa | agagagaaga | aatcttatcg | gagatgaaaa | 660 |
| aggttactga | aggtgttgtc | gatgtcatcg | tctacccaag | cgctgcagat | aaaaccaaaa | 720 |
| accgaggctt | tgccttcgtg | gagtatgaga | gtcatcgagc | agctgccatg | gcgaggagga | 780 |
| aactgctacc | aggaagaatt | cagttatggg | gacatggtat | tgcagtagac | tgggcagagc | 840 |
| cagaagtaga | agttgatgaa | gatacaatgt | cttcagtgaa | aatcctatat | gtaagaaatc | 900 |
| ttatgctgtc | tacctctgaa | gagatgattg | aaaaggaatt | caacaatatc | aaaccaggtg | 960 |
| ctgtggagag | ggtgaagaaa | attcgagact | atgcttttgt | gcacttcagt | aaccgagaag | 1020 |
| atgcagttga | ggctatgaaa | gctttaaatg | gcaaggtgct | ggatggttcc | cccattgaag | 1080 |
| tcaccctagc | aaaaccagtg | gacaaggaca | gttatgttag | gtatacccga | ggcacaggtg | 1140 |
| gaaggggcac | catgctgcaa | ggagagtata | cctactcttt | gggccaagtt | tatgatccca | 1200 |
| ccacaaccta | ccttggagct | cctgtcttct | atgccccca | gacctatgca | gcaattccca | 1260 |
| gtcttcattt | cccagccacc | aaaggacatc | tcagcaacag | agccattatc | cgagcccctt | 1320 |
| ctgttagagg | ggctgcggga | gtgagaggac | tgggcggccg | tggctatttg | gcatacacag | 1380 |
| gcctgggtcg | aggataccag | gtcaaaggag | acaaaagaga | agacaaactc | tatgacattt | 1440 |
| tacctgggat | ggagctcacc | ccaatgaatc | ctgtcacatt | aaaaccccaa | ggaattaaac | 1500 |
| tcgctcccca | gatattagaa | gagatttgtc | agaaaaataa | ctggggacag | ccagtgtacc | 1560 |
| agctgcactc | tgctattgga | caagaccaaa | gacagctatt | cttgtacaaa | ataactattc | 1620 |
| ctgctctagc | cagccagaat | cctgcaatcc | accctttcac | acctccaaag | ctgagtgcct | 1680 |
| ttgtggatga | agcaaagacg | tatgcagccg | aatacaccct | gcagaccctg | ggcatcccca | 1740 |

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<210> 99

<211> 5730

<212> DNA

<213> NM_019008.4| Homo sapiens hypothetical protein FLJ20232 (FLJ20232), mRNA

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| cgtgtcccgg | aagtgaaggg | gccatgttga | tgggtgaccc | ggggagaggt | acccggccag | 120 |
| aggcgagtcc | tgcggagtgg | tagcgcgcac | ggcctgcggt | gtgacaccca | gcccctgcca | 180 |
| gtcccccatg | gccccgtgga | gccgagaggc | ggtgctgagt | ctctatcggg | ctctgttgcg | 240 |
| ccagggccga | cagcttcgct | acactgatcg | agacttctac | tttgcctcca | tccgccgtga | 300 |
| attccgaaaa | aatcagaagc | tagaggacgc | tgaggcccgg | gagaggcagc | tggagaaggg | 360 |
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| 180 | cagccccgca | tccagtcaga | gagcgccggt | gcggggccga | cccgctggga | tgtcagcgcg |
| 240 | aagaagagca | gtaccagggc | gggccatgtc | gcgccagagg | cgggccgagg | ggtcagcggc |
| 300 | aacgatgacc | acggatcatc | tcaaaggtgg | cgactcctca | cacgagtgac | tcccgcacat |
| 360 | ggagagaact | caaacaaata | atggacttat | tacctggagg | tgctgacgtc | aatcccttta |
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| 1200 | ttgcaggtca | ctgtggggac | ccctactggc | tacttgacct | cacgcccgac | cggaccctac |
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| gga | act | act | act | act | ict | .ct | ct | ct | ct | ct | ct | ct | ct | ct | C1 | C | .c | .C | ιC | ac | a | a | a | а | â | | ι | a | a | a | jā | ga | g | Ιg | 36 | g | g | : c | t 9 | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | 1 | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | E | - | - 1 | | | | | (| (| Ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | Ć | Ç |
|-----|-----|-----|-----|-----|-----|------|-----|-----|-----|----|----|----|----|----|----|-----|-----|----|-----|----|----|----|---|---|---|---|---|---|---|----|----|----|----|----|------------|---|------------|------------|-----|---|---|---|---|---|---|---|---|---|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|----|---|----|----|----|----|----|----|----|---|---|---|---|---|---|---|---|---|---|-----|-----|-----|-----|-----|-----|-----|-----|-----|---|-----|-----|-----|-----|-----|-----|-----|-----|-----|---|---|---|---|---|---|-----|-----|-----|-----|-----|-----|-----|---|-----|---|-----|-----|-----|-----|-----|-----|----|----|
| ctt | ggc | ggc | ggc | ggc | jgc | gc | gc | gc | gc | gc | gc | gc | gc | go | g | g | g | ıg | Ιg | gg | g | g | g | õ | ç | | = | 2 | C | C | JC | g | g | :g | <u> </u> | C | c | : 0 | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | 1 | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | = | - | : (| : (| : (| : (| : (| | (| (| C | C | C | C | C | C | C | C | C | C | C | C | C | C | C | C | (| (| (| (| (| (| (| (| (| (| (| (| (| (| (| (| (| (|
| caa | gca | gca | gca | gca | jca | ca | ca | ca | ca | ca | ca | ca | ca | ca | Cā | C | C | IC | JC | gc | g | g | g | õ | ç | | ı | 1 | a | a | ıa | a | a | Įα | jā | g | g | : Q | t (| t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | 1 | 1 | 1 | . 1 | 1 | 1 | 1 | 1 | 1 | 1 | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | _ | - | | | | | : (| | (| ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | Ć | Ć | Ć | Ć | Ć | Ć | Ć | Ć | Ć | Ć | Ć | Ć | Ć | Ć | Ć | Ć | Ć | Ć | Ć | Ć | Ć | • | • |
| gga | cct | cct | cct | cct | ct | ct | ct | ct | ct | ct | ct | ct | ct | C1 | C† | C. | C. | c | c | CC | C | C | C | C | (| | : | 2 | c | C | gc | g | g | ιg | 10 | a | a | Jā | g | g | g | g | g | g | g | g | g | g | g | ç | ć | ć | ć | ć | ć | Ć | Ć | Ć | ļ | Ć | Ć | Ć | Ć | Ć | Ć | ć | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | ç | 9 | g | g | 9 | 9 | 9 | 9 | J | J | J | J | J | J | į | ä | ā | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ē | ć | ē | ć | ć | ć | ć | ć | ć | ä | į |
| gcg | ctc | ctc | ctc | ctc | ctc | :tc | tc | tc | tc | tc | tc | tc | to | t | t | t | :t | :t | :t | ct | C | C. | C | C | (| | : | = | c | C | gc | g | g | :g | 29 | C | C | | C | c | c | C | c | c | c | c | c | C | c | C | C | C | C | C | C | (| (| (| | (| (| (| (| (| (| (| C | C | C | C | C | C | C | C | C | C | C | C | c | c | c | С | c | = | = | : | - | - | - | - | | : (| | (| (| C | C | C | C | C | C | C | C | C | (| (| (| (| (| (| (| (| (| (| (| (| (| (| (| (| (| (| (| (| (| (| • | |
| gga | cgc | cgc | cgc | cgc | :gc | :gc | gc | gc | gc | gc | gc | gc | go | g | g | g | g | g | :g | ΞQ | C | C | c | C | (| |) | 9 | g | .g | ą | a | a | ιa | 16 | a | а | . 6 | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | 1 | 1 | 1 | :1 | 1 | 1 | 1 | 1 | 1 | 1 | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | t | - | | | | | | | | ć | ć | ã | ã | ã | ã | ã | ã | ã | ã | ã | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | |
| gct | cgc | cgc | cgc | cgc | gc | :gc | gc | gc | gc | gc | gc | gc | gc | g | g | g | g | :g | :g | ٥ç | C | C | C | C | (| | j | 9 | g | g | Ę | t | t | :t | 21 | C | C | 30 | g | g | g | g | g | g | g | g | g | g | ō | ē | Ć | Ę | ē | ē | ē | Ç | Ç | Ç | ļ | Ç | Ç | Ç | Ç | Ç | Ç | ē | ē | ē | ē | ē | ē | ē | ē | ē | ē | ē | ē | ē | ç | c | c | g | g | 9 | J | J | J | J | J | J | J | ļ | ļ | • | (| (| (| (| (| (| (| (| (| (| (| (| (| (| (| (| • | • | • | • | • | • | • | (| • | (| • | • | • | • | • | • | • | • |
| cca | gcg | gcg | gcg | gcg | ącg | ıcg | cg | cg | cg | cg | СĈ | cζ | ςį | Сį | C | C | JC: | јС | jς | gc | g | g | g | ç | Ģ | 1 |) | 9 | g | ıg | 36 | g | g | 36 | 36 | g | g | <u> </u> | C | c | C | c | c | c | c | c | c | c | c | c | C | • | (| (| (| | | | | | | | | | | (| (| C | (| (| (| (| (| (| (| C | C | C | c | c | c | c | С | = | = | = | - | - | - | - | - | | | Ç | Ç | Ç | Ç | Ç | Ç | Ç | Ç | Ç | Ç | Ç | Ç | Ç | Ç | Ç | Ç | Ç | Ç | Ç | Ç | Ç | Ç | Ç | Ç | ę | Ç | ę | Ç | Ç | Ç | Ç | Ç | Ç | • | |
| ato | gct | gct | gct | gct | gct | jct | ct | ct | ct | ct | ct | ct | C1 | C1 | C. | IC. | JC. | ĵС | jς | gc | g | g | g | ç | Ç | L | a | a | a | :a | 2 | C | C | :c | C (| C | C | C (| t | t | t | t | t | t | t | t | t | t | t | t | :1 | :1 | :1 | :1 | :1 | :1 | :1 | :1 | | :1 | :1 | :1 | :1 | :1 | :1 | 1 | 1 | t | 1 | 1 | 1 | 1 | 1 | 1 | 1 | t | t | t | t | t | t | t | t | t | t | _ | - | - | - | - | - | | : (| | • | (| (| (| (| (| (| (| (| (| • | (| (| (| (| (| • | • | • | • | • | • | • | • | • | • | • | • | • | • | • | • | • | |
| gga | cga | cga | cga | cga | cga | :ga | :ga | ga | ga | ga | ga | ga | ga | ga | g | :g | :g | :g | Ξg | CQ | C | C | c | C | (| : | = | С | c | :c | t | t | t | :t | t1 | t | t | _ 1 | t | t | t | t | t | t | t | t | t | t | t | t | . 1 | : 1 | 1 | 1 | 1 | : 1 | : 1 | : 1 | | : 1 | : 1 | : 1 | : 1 | : 1 | : 1 | 1 | 1 | t | 1 | 1 | 1 | 1 | 1 | 1 | 1 | t | t | t | t | t | t | t | t | t | t | = | - | - | - | - | - | - | | | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | - | | |
| tgt | acc | aco | acc | acc | acc | ıcc' | ıcc | .CC | .cc | cc | cc | c | C | C | .C | ιC | ιC | ιC | a C | ac | a | a | a | ā | i | L | a | a | a | :a | tá | t | t | :t | C† | C | : C | = | c | C | c | C | c | c | c | c | c | C | . C | . (| :(| : (| | | | : (| : (| : (| | : (| : (| : (| : (| : (| : (| (| (| (| (| (| (| (| (| (| (| (| (| (| C | C | C | c | c | C | _ | _ | - | - | - | - | - | - | | : (| | (| (| (| (| (| (| (| (| (| | • | • | • | • | • | | | | | | | | | | | | | | | | | | |
| ago | tcg | tcg | tcg | tcg | tcg | :cg | :cg | cg | cg | СĆ | СĆ | ςį | С | C | C | :c | :c | :c | tc | t | t | t | t | t | | L | a | a | a | ιa | aa | a | .a | 12 | a | a | ıa | 1 | a | a | а | а | а | а | а | а | а | а | ā | ā | ιē | ιź | lá | lá | lá | lá | lá | lá | ı | lá | lá | lá | lá | lá | lá | . 6 | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ē | ĉ | ĉ | а | а | a | a | a | ı | ı | ı | ı | ı | l | l | li | | ě | ě | ě | ě | ě | ě | ě | ě | ě | | | | | | | . : | . : | . : | . : | . : | . : | . : | | . : | | . : | . : | . : | . : | . : | . : | | Į, |
| aat | ctg | cto | ctg | ctg | ctg | :tg | :tg | tg | tg | tς | ŧς | ŧς | t | :t | :t | :t | :t | :t | ct | C1 | C. | C | c | (| (| ι | a | a | a | :a | ta | t | :t | Ξt | t. | t | ιt | a . | a | а | а | а | ĉ | ĉ | а | а | а | 2 | ē | ā | : 7 | : 7 | : 6 | : 6 | : 6 | : 6 | : 6 | : 6 | : | : 6 | : 6 | : 6 | : 6 | : 6 | : 6 | | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ć | ã | ã | ĉ | а | a | a | a | ı | ı | ı | ı | ı | ı | ľ | ı. | Ľ | . 1 | . 1 | . 1 | . 1 | . 1 | . 1 | . 1 | . 1 | . 1 | Ľ | | | | | | Ľ | Ľ | Ľ | Ľ | Ľ | Ľ | Ľ | | Ľ | | Ľ | Ľ | Ľ | Ľ | Ľ | Ľ | ľ. | ľ |
| tt1 | tct | tct | tct | tct | tct | :ct | :ct | ct | ct | ct | ct | C1 | C | c. | :c | :c | :c | :c | tc | t | t | t | t | 1 | | ι | a | a | a | :a | ta | t | :t | Εt | t: | t | ļτ | 3. | g | g | ç | ç | ē | ē | ē | ē | ē | ē | Ç | Į | ļ | Į | ļ | ļ | ļ | J | J | J | J | J | J | J | J | J | J | ! | 9 | ç | 9 | 9 | 9 | 9 | 9 | 9 | 9 | ç | ç | ç | ć | ç | ç | ç | ç | 9 | 9 | J | J | J | J | J | J | J | J. | Į. | ŀ | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | ŀ | • | • | • | • | • | ı. | | ı. | | ı. | ı. | ı. | ı. | ı. | ı. | Į. | J |
| aa1 | aat | aat | aat | aat | aat | ıat | ıat | at | .at | at | a۱ | a٦ | a' | a | ıa | ιa | ıa | ıa | aa | aa | a | a | a | ä | i | J | 9 | g | g | ĵĈ | 99 | g | ιg | 10 | a | a | :2 | C | c | C | C | C | C | C | C | C | C | C | : (| : (| : (| : (| : (| : (| : (| | | | - | | | | | | | : (| • | (| • | • | • | • | • | • | • | (| (| (| (| C | C | C | C | c | C | C | = | = | = | = | = | 2 | 2 | : | : | | | | | | | | | | : | : | : | : | : | : | : | : | : | : | : | : | : | | : | | : | : | : | : | : | : | | : |
| ato | ttt | ttt | ttt | ttt | ttt | :tt | :tt | tt | .tt | tt | t۱ | t1 | t | :t | :t | :t | :t | Ξŧ | tt | t۱ | t | t | t | 1 | • | - | t | t | t | :t | t | t | :t | t٦ | t. | t | :t | Ξ. | c | C | c | C | C | C | C | C | C | C | : 0 | :0 | :(| -(| : (| : (| : (| - (| - (| - (| - | - (| - (| - (| - (| - (| - (| | | (| | | | | | | | (| (| (| (| C | C | C | C | c | C | = | _ | _ | _ | _ | _ | 2 | - | | : | | | | | | | | | | : | | | | | | | | | | | | | | | | | | | | | | | : |
| tta | tat | tat | tat | tat | tat | at | :at | at | .at | at | at | a1 | a | :a | :a | :a | :a | :a | ta | ta | t | t | t | 1 | • | J | g | g | g | ıç | a | a | ιa | 12 | a | a | ιā | a | a | а | a | ĉ | ã | ã | ĉ | ĉ | ĉ | ē | į | Įā | Į | Į | Į | Į | Į | J | J | J | J | J | J | J | J | J | J | į | • | č | • | • | • | • | • | • | • | č | č | č | ā | ć | ć | ã | ĉ | а | a | a | a | a | a | a | a | ı | ı | l | l | li | l | L | L | L | L | L | ι | ι | ι | ι | ι | ι | ι | L | ι | L | ι | ι | ι | ι | ι | ι | ι | ι |
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | l | a | a | ıa | aa | a | a | lā | a | a | a | ā | ā | ā | ā | ā | ā | ā | ā | ιā | ιā | lä | lä | ı | ı | ı | ı | ı | ı | ı | ı | ı | ı | ı | ı | ı | l | lě | ě | lě | ě | ě | ě | ć | 2 | 2 | ē | ã | а | а | a | a | a | a | a | 1 | ı | ı | l | t | Li | t | L | L | L | L | L | l | l | l | l | l | l | l | L | l | L | l | l | l | l | l | l | ι | ι |
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |

<210> 110

<211> 3 2 6 2

<212> DNA

 $<\!\!213\!\!>$ NM_002860.2| Homo sapiens aldehyde dehydrogenase 18 family, member A1 (PYCS/ALDH18A1), mRNA

| <400> 110 agggggcgaa | ggcggcggtg | gtgaggaaga | tactttggtt | agtgaccaca | tcgcagcatg | 60 |
|-------------------------|------------|------------|------------|------------|------------|-----|
| ttgagtcaag | tttaccgctg | tgggttccag | cccttcaacc | aacatcttct | gccctgggtc | 120 |
| aagtgtacaa | ccgtcttcag | atctcattgt | atccagcctt | cagtcatcag | acatgttcgt | 180 |
| tcttggagca | acatcccgtt | tatcactgta | cccctcagtc | gtacacatgg | aaagtccttc | 240 |
| gcccaccgca | gtgagctgaa | gcatgccaag | agaatcgtgg | tgaagctcgg | cagtgccgtg | 300 |
| gtgacccgag | gggatgaatg | tggcctggcc | ctggggcgct | tggcatctat | tgttgagcag | 360 |
| gtatcagtgc | tgcagaatca | gggcagagag | atgatgctgg | tgaccagtgg | agccgtagcc | 420 |
| tttggcaaac | aacgcttgcg | ccatgagatc | cttctgtctc | agagcgtgcg | gcaggccctc | 480 |
| cactcggggc | agaaccagct | gaaagaaatg | gcaattccag | tcttagaggc | acgagcctgt | 540 |

| gcagctgccg | gacagagtgg | gctgatggcc | ttgtatgagg | ctatgtttac | ccagtacagc | 600 |
|---------------------|------------|------------|------------|------------|------------|------|
| atctgtgctg | cccagatttt | ggtgaccaat | ttggatttcc | atgatgagca | gaagcgccgg | 660 |
| aacctcaatg | gaacacttca | tgaactcctt | agaatgaaca | ttgtccccat | tgtcaacaca | 720 |
| aatgatgctg | ttgtccccc | agctgagccc | aacagtgacc | tgcagggggt | aaatgttatt | 780 |
| agtgttaaag | ataatgatag | cctggctgcc | cgactggctg | tggaaatgaa | aactgatctc | 840 |
| ttgattgttc | tttcagatgt | agaaggcctt | tttgacagcc | ccccaggttc | agatgatgca | 900 |
| aagcttattg | atatattta | tcccggagat | cagcagtctg | tgacatttgg | aaccaagtct | 960 |
| agagtgggaa | tgggtggcat | ggaagccaag | gtgaaagcag | ccctctgggc | tttgcaaggt | 1020 |
| ggcacttctg | ttgttattgc | caatggaacc | cacccaaagg | tgtctgggca | cgtcatcaca | 1080 |
| gacattgtgg | aggggaagaa | agttggtacc | ttcttttcag | aagtaaagcc | tgcaggccct | 1140 |
| actgttgagc | agcagggaga | aatggcgcga | tctggaggaa | ggatgttggc | caccttggaa | 1200 |
| cctgagcaga | gagcagaaat | tatccatcat | ctggctgatc | tgttgacgga | ccagcgtgat | 1260 |
| gagatcctgt | tagccaacaa | aaaagacttg | gaggaggcag | aggggagact | tgcagctcct | 1320 |
| ctgctgaaac | gtttaagcct | ctccacatcc | aaattgaaca | gcctggccat | cggtctgcga | 1380 |
| cagatcgcag | cctcctccca | ggacagcgtg | ggacgtgttt | tgcgccgcac | ccgaatcgcc | 1440 |
| aaaaacttgg | aactggaaca | agtgactgtc | ccaattggag | ttctgctggt | gatctttgaa | 1500 |
| tctcgtcctg | actgtctacc | ccaggtggca | gctttggcta | tcgcaagtgg | caatggcttg | 1560 |
| ttactcaaag | gagggaagga | ggctgcacac | agcaaccgga | ttctccacct | cctgacccag | 1620 |
| gaggctctct | caatccatgg | agtcaaggag | gccgtgcaac | tggtgaatac | cagagaagaa | 1680 |
| gttgaagatc | tttgccgcct | agacaaaatg | atagatctga | tcattccacg | tggctcttcc | 1740 |
| cagctggtca | gagacatcca | gaaagctgct | aaggggattc | cagtgatggg | gcacagcgaa | 1800 |
| gggatctgtc | acatgtatgt | ggattccgag | gccagtgttg | ataaggtcac | caggctagtc | 1860 |
| agagact c ta | aatgtgaata | tccagctgcc | tgtaatgctt | tggagacttt | gttaatccac | 1920 |
| cgggatctgc | tcaggacacc | attatttgac | cagatcattg | atatgctgag | agtggaacag | 1980 |
| gtaaaaattc | atgcaggccc | caaatttgcc | tcctatctga | ccttcagccc | ctccgaagtg | 2040 |
| aagtcactcc | gaactgagta | tggggacctg | gaattatgca | ttgaagtagt | ggacaacgtt | 2100 |
| caggatgcca | ttgaccacat | ccacaagtat | ggcagctccc | acacggatgt | catcgtcaca | 2160 |
| gaggacgaaa | acacagcgga | gttcttcctg | cagcacgtag | acagtgcctg | tgtgttctgg | 2220 |
| aatgccagca | ctcgcttttc | tgatggttac | cgctttggac | tgggagctga | agtgggaatc | 2280 |
| agtacatcga | gaatccacgc | ccggggacca | gtaggacttg | agggactgct | tactactaag | 2340 |
| tggctgctgc | gagggaagga | ccacgtggtc | tcagatttct | cagagcatgg | aagtttaaaa | 2400 |
| tatcttcatg | agaacctccc | tattcctcag | agaaacacca | actgaaaaga | gccaggaaaa | 2460 |
| cccgggaatt | ttccaaaagg | tcttcacgtt | aaacttgtct | tatctcagga | gagagcccgc | 2520 |
| tcttgtctcc | cagttcctgg | tagggtctgc | ctgttggaaa | gtgtacctgg | atgcttctgg | 2580 |
| | | | | | | |

```
2640
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tttttttaaa gtaagaaaat agttgctacc gatagggact ttgccaagtc caattatctt
                                                                    2700
                                                                    2760
ctaggattga aaggtgcatt ttccccataa aaaaggcgag gaaaacccat ggctgctttg
tgtcacctca gtgacttaca gtcccccttc gcatttagtt ggtactagag ccagtcatcc
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                                                                    2880
ttggaatttt agaaaagggg caactcttct ttttagcatt ctcatcagaa agtcacaaaa
                                                                    2940
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                                                                    3000
                                                                    3060
aataagcatt ccagatgctt gcagcttcct gcatccagga gatgctgtgt cccccgtgat
gcagctggaa cccaagctgc agcaggagat gcaagtttca ggatgttccc cactgagctg
                                                                    3120
gaggaatatc tacagcagtg atgcttgaaa tttttgtatg aattattttg tcgtcctacc
                                                                    3180
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                                                                     3240
                                                                     3262
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<210> 111

<211> 2899

<212> DNA

<213> NM_005655.1 \mid Homo sapiens TGFB inducible early growth response (TIEG), mRNA

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| accatgctca | acttcggtgc | ctctctccag | cagactgcgg | aggaaagaat | ggaaatgatt | 180 |
| tctgaaaggc | caaaagagag | tatgtattcc | tggaacaaaa | ctgcagagaa | aagtgatttt | 240 |
| gaagctgtag | aagcacttat | gtcaatgagc | tgcagttgga | agtctgattt | taagaaatac | . 300 |
| gttgaaaaca | gacctgttac | accagtatct | gatttgtcag | aggaagagaa | tctgcttccg | 360 |
| ggaacacctg | attttcatac | aatcccagca | ttttgtttga | ctccacctta | cagtccttct | 420 |
| gactttgaac | cctctcaagt | gtcaaatctg | atggcaccag | cgccatctac | tgtacacttc | 480 |
| aagtcactct | cagatactgc | caaacctcac | attgccgcac | ctttcaaaga | ggaagaaaag | 540 |
| agcccagtat | ctgccccaa | actccccaaa | gctcaggcaa | caagtgtgat | tcgtcataca | 600 |
| gctgatgccc | agctatgtaa | ccaccagacc | tgcccaatga | aagcagccag | catcctcaac | 660 |
| tatcagaaca | attcttttag | aagaagaacc | cacctaaatg | ttgaggctgc | aagaaagaac | 720 |
| ataccatgtg | ccgctgtgtc | accaaacaga | tccaaatgtg | agagaaacac | agtggcagat | 780 |
| gttgatgaga | aagcaagtgc | tgcactttat | gacttttctg | tgccttcctc | agagacggtc | 840 |
| atctgcaggt | ctcagccagc | ccctgtgtcc | ccacaacaga | agtcagtgtt | ggtctctcca | 900 |

| cctgcagtat | ctgcaggggg | agtgccacct | atgccggtca | tctgccagat | ggttcccctt | 960 |
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| cctgccaaca | accctgttgt | gacaacagtc | gttcccagca | ctcctcccag | ccagccacca | 1020 |
| gccgtttgcc | cccctgttgt | gttcatgggc | acacaagtcc | ccaaaggcgc | tgtcatgttt | 1080 |
| gtggtacccc | agcccgttgt | gcagagttca | aagcctccgg | tggtgagccc | gaatggcacc | 1140 |
| agactctctc | ccattgcccc | tgctcctggg | ttttcccctt | cagcagcaaa | agtcactcct | 1200 |
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| tactttaaaa | gttcccatct | gaaggcccac | acgaggacgc | acacaggaga | aaagcctttc | 1320 |
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| aggcgaaccc | acacgggtga | gaagaaattt | gcgtgcccca | tgtgtgaccg | gcggttcatg | 1440 |
| aggagtgacc | atttgaccaa | gcatgcccgg | cgccatctat | cagccaagaa | gctaccaaac | 1500 |
| tggcagatgg | aagtgagcaa | gctaaatgac | attgctctac | ctccaacccc | tgctcccaca | 1560 |
| cagtgacaga | ccggaaagtg | aagagtcaga | actaactttg | gtctcagcgg | gagccagtgg | 1620 |
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| ccccacagcc | tggcacgaag | gccccgcctg | ggttaggtga | ctaaaagggc | ttcggccaca | 1740 |
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| attcctttga | atatttttg | aaggtttcag | atgaggtcaa | cacaggtagc | acagattttg | 1860 |
| aatctgtgtg | catatttgtt | actttacttt | tgctgtttat | acttgagacc | aacttttcaa | 1920 |
| tgtgattctt | ctaaagcact | ggtttcaaga | atatggaggc | tggaaggaaa | taaacattac | 1980 |
| ggtacagaca | tggagatgta | aaatgagttt | gtattattac | aaatattgtc | atctttttct | 2040 |
| agagttatct | tctttattat | t c ctagtctt | tccagtcaac | atcgtggatg | tagtgattaa | 2100 |
| atatatctag | aactatcatt | tttacactat | tgtgaatatt | tggaattgaa | cgactgtata | 2160 |
| ttgctaagag | ggcccaaaga | attggaatcc | tccttaattt | aattgctttg | aagcatagct | 2220 |
| acaatttgtt | tttgcatttt | tgttttgaaa | gtttaacaaa | tgactgtatc | taggcatttc | 2280 |
| attatgcttt | gaactttagt | ttgcctgcag | tttcttgtgt | agatttgaaa | attgtatacc | 2340 |
| aatgtgtttt | ctgtagactc | taagatacac | tgcactttgt | ttagaaaaaa | aactgaagat | 2400 |
| gaaatatata | ttgtaaagaa | gggatattaa | gaatcttaga | taacttcttg | aaaaagatgg | 2460 |
| cttatgtcat | cagtaaagta | cctttatgtt | atgaggatat | aatgtgtgct | ttattgaatt | 2520 |
| agaaaattag | tgaccattat | tcacaggtgg | acaaatgttg | tcctgttaat | ttataggagt | 2580 |
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| cattgcttaa | aatatatagt | gaaaaatgtc | actatatctt | cccatttaac | attgtttttg | 2760 |
| tatattgggt | gtagatttct | gacatcaaaa | cttggaccct | tggaaaacaa | aagttttaat | 2820 |
| taaaaaaaat | ccttgtgact | tacaatttgc | acaatatttc | ttttgttgta | ctttatatct | 2880 |
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tgtttacaat aaagaattc 2899

<210> 112

<211> 3138

<212> DNA

 $<\!\!213\!\!>$ NM_018223.1| Homo sapiens checkpoint with forkhead and ring finger domains (CHFR), mRNA

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| | | agggcgagcc | | | | 180 |
| | | gcgacctttc | | | | 240 |
| | | atgaaaaatc | | | | 300 |
| | | agctgaaggt | | | | 360 |
| | | | | | | 420 |
| | | tgtacaggaa | | | | 480 |
| | | agcaaggcat | | | | 540 |
| | | taaaactcct | | | | 600 |
| | | ggtctggggg | | | | |
| | | tctccagctt | | | | 660 |
| | | cccaggatca | | | | 720 |
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| tgcccaaacc | gtccacgagg | acgtcagagc | agcggctggg | aagccagaca | agatggagga | 840 |
| gacgctgaca | tgcatcatct | gccaggacct | gctgcacgac | tgcgtgagtt | tgcagccctg | 900 |
| catgcacacg | ttctgcgcgg | cttgctactc | gggctggatg | gagcgctcgt | ccctgtgtcc | 960 |
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| | | gctgcttcca | | | | 1440 |
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<211> 2466

<212> DNA

<213> NM_024645.1| Homo sapiens hypothetical protein FLJ13842 (FLJ13842), mRNA

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| gcccctctgc | ctcgcggaaa | agcctgatga | agtcctccga | tattgatcag | gatttattca | 180 |
| cagacagtta | ctgcaaggtg | tgcagtgcac | agctgatctc | cgaatcgcag | cgtgtggccc | 240 |
| actacgagag | tcgaaaacat | gcaagcaaag | tccgactgta | ttacatgctt | caccccaggg | 300 |
| atggagggtg | tcctgccaag | aggctccggt | cagaaaatgg | aagtgatgcc | gacatggtgg | 360 |
| ataagaacaa | gtgctgcaca | ctctgcaaca | tgtcattcac | ttcagcggtg | gtggccgatt | 420 |
| cccattatca | aggcaaaatc | cacgccaaga | ggttaaaact | cttgctagga | gagaagaccc | 480 |
| cattaaagac | cacagcaaca | cccctgagcc | cacttaagcc | cccacggatg | gacactgctc | 540 |
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| gaactgtggg | ctttaattgg | gggatactga | tcgtttggaa | agaagtgaga | aaattctgaa | 1740 |
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| actgctgagc | caagactcag | tcactctgga | aagagcatga | ccgataaaga | aaacagttcc | 1860 |
| tttctgatgg | ggagcgtctg | agtgcagatc | atgaggctct | ttctctaggt | ttaattcttt | 1920 |
| tccatggtga | ccggacttgg | tgtcttgtag | cctggttacg | aagtgggacg | ttgagcttct | 1980 |
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<210> 114

<211> 3658

<212> DNA

<213> NM_025195.2| Homo sapiens tribbles homolog 1 (Drosophila) (TRIB1), mRNA

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| gcgccgcgaa | aaagttcccc | cggcttttgc | tggagactca | tcgttttggg | aagtgcattt | 180 |
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| aagcttagga | agttcgtctt | ctccacggag | gagagaaccc | agcttagact | agaaagtcta | 1260 |
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| catcaggatg | aaagctgctg | aactcggcat | ggcgcctcct | cttctctgtt | gggatgagtg | 1860 |
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| aatatactgt | attttgagaa | atggcacaaa | aacaggcagt | catctttaag | ggctatgcct | 3180 |
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| 180 | cgggctgggg | cacctggggg | cggctgcggg | ccgcgccagc | gtcgcctcgg | ctgcagcccc |
| 240 | gcggcggccg | cgccggtcct | agggctgcgg | cgctgtagcg | cggcaggagg | gcgccggccg |
| 300 | gcgcgaccgc | tccggctcct | gggctcctcc | gctgtgggcc | cggggcaggc | cgggaggcag |
| 360 | cccgccgccc | ccgctctgcg | ccccgcagcg | cgcccgccgt | gctctgccgg | ctcccgccgg |
| 420 | ggggccatgg | gcgggcgcgc | cggcgggcgc | gcgggagcct | cgcggggctg | cgagcgcccg |
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| 1020 | aagaaggcaa | tcatgcagta | ttgactgttt | attacacttc | caatttctac | atggaagttg |
| 1080 | cactatgaaa | tgaatatgaa | ttaaccttga | ttcaactcat | cttccttaat | tgcagtatgg |
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<211> 3919

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<211> 7401

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| ctctgcctag | cagcagtgct | gccacagccc | tgctgatcca | gcgcaccaat | gaggaggaga | 1560 |
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| ccgagttccg | gcagcaggtg | ctggagcagc | acctggatat | gggccggccc | ccggtgccgg | 1680 |
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| gtggccgtga | ccctccgcag | cactttgggg | ggccaccacc | tctgatttcg | cccaagcccc | 1800 |

| agctccatgc | tgcacccacg | gccctctgga | accccgtgtc | cctgatggac | aacaccttgg | 1860 |
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| agacgcggcg | ggccgaaagc | cactctctgc | acagccaccc | ggctgcattt | gagcccagcc | 1920 |
| gccaggcagc | cgtgccgctg | gtgaaggtgg | agcgggtctt | ctgcccggag | aaagcagagg | 1980 |
| aggggccacg | gaagcgtgag | cctgcccctc | tggacaagta | ccagccacct | ccgccgccac | 2040 |
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| aaaggttggg | | | | | | 4260 |
| attttcttca | | | | | | 4320 |
| aaaagctctt | | | | | | 4380 |
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| | | | | attaaaatat | | 4740 |
| | | | | tggcgcctat | | 4800 |
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<210> 118

<211> 2745

<212> DNA

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| ggagcccgcc | ggtgaggccg | ggccacgctc | agacacttcg | atcgtcgagt | ctgtcactgg | 180 |
| gcatggcggg | tcagttccgc | agctacgtgt | gggacccgct | gctgatcctg | tcgcagatcg | 240 |
| tcctcatgca | gaccgtgtat | tacggctcgc | tgggcctgtg | gctggcgctg | gtggacgggc | 300 |
| tagtgcgaag | cagcccctcg | ctggaccaga | tgtttgacgc | cgagatcctg | ggcttttcca | 360 |
| ccctccagg | ccggctctcc | atgatgtcct | tcatcctcaa | cgccctcacc | tgtgccctgg | 420 |
| gcttgctgta | cttcatccgg | cgaggaaagc | agtgtctgga | tttcactgtc | actgtccatt | 480 |
| tctttcacct | cctgggctgc | tggttctaca | gctcccgttt | cccctcggcg | ctgacctggt | 540 |
| ggctggtcca | agccgtgtgc | attgcactca | tggctgtcat | cggggagtac | ctgtgcatgc | 600 |
| ggacggagct | caaggagata | cccctcaact | cagcccctaa | atccaatgtc | tagaatcagg | 660 |
| ccctttggac | atcctgctga | cacttgggcc | ccttaacacc | ttgggctgct | cagaccctcc | 720 |
| agatgaggtc | cagcccagat | ctgagaggaa | ccctggaaat | gtgaagtctc | tgttggtttg | 780 |
| ggagagatag | tgagggcctg | tcaaagaagg | caggtagcag | tcagcatgac | agctgcaaga | 840 |
| atgacctctg | tctgttgaag | ccttggtatc | tgagaggtca | ggaaggggac | ctctttgagg | 900 |
| gtaataacag | aattggaacc | atgccactct | tgagccacaa | tacctgtcac | cagcctgttg | 960 |
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| acggttgaga | ttgagagaga | tcagcgcagc | caggcaaggg | aactttaaag | aattattagg | 1740 |
| ccaccttctc | cctttcctgg | accccagagt | cattcctcca | tttggttaaa | atactcagtg | 1800 |
| cagggaactc | ttacatcctg | tctccttcac | ttgcagcgtc | ccctgctatg | cctcaggtga | 1860 |
| accacataat | tcttgggttt | ccgttcctac | : ttgctagtga | ı tttctgaaca | tgttcaatgg | 1920 |
| agcggcacac | agtctagacc | cacttccgca | ttgaaacctt | cactgttcct | ctttggtttc | 1980 |
| ttcagagctt | tcccaagaga | gctgtcagtt | : ttcagctgtc | agtaacacaa | atgagtttat | 2040 |
| | | | | | | |

| ggtaacacaa | atgagttttg | ctatctctct | gagaagctca | tctgacctcc | tgactctcag | 2100 |
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| ccctacagag | tagggagttg | atgctgacag | gatgaagatt | taggaataaa | tatgcctggg | 2160 |
| aagagactgg | gaaggttcta | gggtgaggca | cctcagtaac | tcatggtacc | ttggccaagt | 2220 |
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| aaagtaacaa | taaaagcagc | ttttagagtt | gagttccaga | gagggcaggg | caatggcagt | 2700 |
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<211> 2152

<212> DNA

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| tggattttct | ctacaaaaat | tattgagcaa | ccctaattaa | cctgattttt | tgctgataat | 180 |
| cactctcaat | ggaatcaaat | cacaaatccg | gggatggatt | gagcggcact | cagaaggaag | 240 |
| cagccctccg | cgcactggtc | cagcgcacag | gatatagctt | ggtccaggaa | aatggacaaa | 300 |
| gaaaatatgg | tggccctcca | cctggttggg | atgctgcacc | ccctgaaagg | ggctgtgaaa | 360 |
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| aaatcggtaa | aatttatgaa | atgagaatga | tgatggattt | taatggcaac | aatagaggat | 480 |
| atgcatttgt | aacattttca | aataaagtgg | aagccaagaa | tgcaatcaag | caacttaata | 540 |
| attatgaaat | tagaaatggg | cgcctcttag | gggtttgtgc | cagtgtggac | aactgccgat | 600 |
| tatttgttgg | gggcatccca | aaaaccaaaa | agagagaaga | aatcttatcg | gagatgaaaa | 660 |
| aggttactga | aggtgttgtc | gatgtcatcg | tctacccaag | cgctgcagat | aaaaccaaaa | 720 |
| accgaggctt | tgccttcgtg | gagtatgaga | gtcatcgagc | agctgccatg | gcgaggagga | 780 |
| aactgctacc | aggaagaatt | cagttatggg | gacatggtat | tgcagtagac | tgggcagagc | 840 |
| cagaagtaga | agttgatgaa | gatacaatgt | cttcagtgaa | aatcctatat | gtaagaaatc | 900 |

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2152
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<211> 3010

<212> DNA

<213> NM_145343.1 \mid Homo sapiens apolipoprotein L, 1 (APOL1), transcript variant 2, mRNA

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| tccctaatgg | gaaacgtggc | tgagacaggg | gagtgagaag | ggtgcgttgc | agaatggtgc | 240 |
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| aggccctgca | gcgacatgga | gggagctgct | ttgctgagag | tctctgtcct | ctgcatctgg | 360 |

| | | | | | | 420 |
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| gctgctggca | | | | | | 540 |
| aaggaaaaag ` | | | | | | 600 |
| | | | aatgaggcag | | | 660 |
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| tacagaaact | ggtttctgaa | agagtttcct | cggttgaaaa | gtgagcttga | ggataacata | 780 |
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| aatgtggtgt | ctggctctct | cagcatttcc | tctggcatcc | tgaccctcgt | cggcatgggt | 900 |
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| acggatgtgg | cccctgtaag | cttctttctt | gtgctggatg | tagtctacct | cgtgtacgaa | 1380 |
| tcaaagcact | tacatgaggg | ggcaaagtca | gagacagctg | aggagctgaa | gaaggtggct | 1440 |
| caggagctgg | aggagaagct | aaacattctc | aacaataatt | ataagattct | gcaggcggac | 1500 |
| caagaactgt | gaccacaggg | cagggcagcc | accaggagag | atatgcctgg | caggggccag | 1560 |
| gacaaaatgc | aaacttttt | ttttttctga | gacagagtct | tgctctgtcg | ccaagttgga | 1620 |
| gtgcaatggt | gcgatctcag | ctcactgcaa | gctctgcctc | : ccgtgttcaa | gcgattctcc | 1680 |
| tgccttggcc | tcccaagtag | ctgggactac | : aggcgcctac | caccatgccc | agctaatttt | 1740 |
| tgtattttta | atagagatgg | ggtttcacca | tgttggccag | gatggtctcg | atctcctgac | 1800 |
| | | | | | agccatcgct | 1860 |
| | | | | | agtttatgga | 1920 |
| actgagtgtt | agggactttg | gcatttccat | agctgagca | agcaggggag | gggttaatgc | 1980 |
| agatggcagt | gcagcaagga | ı gaaggcagga | a acattggago | ctgcaataag | g ggaaaaatgg | 2040 |
| gaactggaga | gtgtggggaa | tgggaagaag | g cagtttactt | t tagactaaag | g aatatattgg | 2100 |
| ggggccgggt | gtagtggct | atgcctgtaa | a tccgagcac | t ttgggaggco | aaggcgggcg | 2160 |
| | | | | | cgtctctact | 2220 |
| | | | | | g ctaactgggc | 2280 |
| | | | | | g ccgagatatc | 2340 |
| | | | | | | |

| gccactgcac | tccagcctgg | gtgacagagc | gagactccat | ctcaaaaaaa | aaaaaaaaa | 2400 |
|------------|------------|------------|------------|------------|------------|------|
| gaatatattg | acggaagaat | agagaggagg | cttgaaggaa | ccagcaatga | gaaggccagg | 2460 |
| aaaagaaaga | gctgaaaatg | gagaaagccc | aagagttaga | acagttggat | acaggagaag | 2520 |
| aaacagcggc | tccactacag | acccagcccc | aggttcaatg | tcctccgaag | aatgaagtct | 2580 |
| ttccctggtg | atggtcccct | gccctgtctt | tccagcatcc | actctccctt | gtcctcctgg | 2640 |
| gggcatatct | cagtcaggca | gcggcttcct | gatgatggtc | attggggtgg | ttgtcatgtg | 2700 |
| atgggtcccc | tccaggttac | taaagggtgc | atgtcccctg | cttgaacact | gaagggcagg | 2760 |
| tggtgggcca | tggccatggt | ccccagctga | ggagcaggtg | tccctgagaa | cccaaacttc | 2820 |
| ccagagagta | tgtgagaacc | aaccaatgaa | aacagtccca | tcgctcttac | ccggtaagta | 2880 |
| aacagtcaga | aaattagcat | gaaagcagtt | tagcattggg | aggaagctca | gatctctaga | 2940 |
| gctgtcttgt | cgccgcccag | gattgacctg | tgtgtaagtc | ccaataaact | cacctactca | 3000 |
| tcaagctgga | | | | | | 3010 |
| | | | | | | |

<210> 121

<211> 2759

<212> DNA

<213> NM_080796.1| Homo sapiens death associated transcription factor 1 (DATF1), transcript variant 2, mRNA $\,$

| <400> 121 tctagccccg | cgccatctcg | gtggccgtcc | gcccactccg | cggcgttcgg | ggaaatggct | 60 |
|----------------------|------------|------------|------------|------------|------------|-----|
| gcgagaccct | agaggcctgc | ggcctgcgga | gcttactcca | cgggaacagc | ctctagataa | 120 |
| tctgagttgt | tgaaaatacg | aagcctgtta | ctcgtgaaca | gtggctgaca | acagtgttgt | 180 |
| tgtgagcctg | gctgtctgct | tggacccaga | ggtttcgtct | gccagggttt | ttggttgtat | 240 |
| ttaggatttc | agggaaaagt | gtccaagctt | tcagtgttgg | agcaggtatg | gacgacaaag | 300 |
| gcgacccgag | caatgaggag | gcacctaagg | ccatcaaacc | caccagcaaa | gagttcagga | 360 |
| aaacatgggg | ttttcgaagg | accactatcg | ccaagcgaga | gggcgcaggg | gacgcggagg | 420 |
| ctgacccact | ggagccgcca | ccccacagc | agcagctggg | cctgtccctg | cggcgcagtg | 480 |
| ggaggcagcc | caagcgcact | gagcgcgtgg | agcagttcct | gaccattgcg | cggcgccgcg | 540 |
| gcaggaggag | catgcctgtc | tccctggagg | attctggtga | gcccacgtcc | tgccccgcca | 600 |
| cagacgccga | gacagcctcc | gagggcagcg | tggaaagcgc | ttctgagacc | agaagcggcc | 660 |
| cccagtctgc | ttccacagct | gtgaaggaac | gaccagcctc | ttctgaaaag | gtgaaaggag | 720 |
| gggatgacca | cgatgacacc | tccgatagtg | acagcgatgg | cctgaccttg | aaagagcttc | 780 |
| agaatcgcct | tcgcaggaag | cgggaacagg | agcccactga | gaggcccctg | aaagggatcc | 840 |
| agagtcgcct | gcggaagaag | cgccgggagg | agggtcccgc | cgagactgtg | ggctccgagg | 900 |

| ccagtgacac | tgtggagggc | gtcctgccca | gtaagcagga | gcccgagaac | gatcaggggg | 960 |
|------------|--------------|--------------|--------------|------------|------------|------|
| ttgtgtccca | ggctgggaaa | gatgacagag | agagtaagtt | ggagggaaag | gcggctcagg | 1020 |
| acatcaaaga | tgaggagcct | ggagacttgg | gccgaccgaa | gcctgaatgt | gagggttacg | 1080 |
| accccaacgc | cctgtattgc | atttgccgcc | agcctcacaa | caacaggttt | atgatttgct | 1140 |
| gtgaccgctg | tgaagaatgg | tttcatggcg | attgtgtggg | catttctgag | gctcgaggga | 1200 |
| ggcttttgga | aaggaatggg | gaagactata | tctgcccaaa | ctgcaccatt | ctgcaagtgc | 1260 |
| aggatgagac | tcattcagaa | acggcagatc | agcaggaagc | taaatggaga | cctggagatg | 1320 |
| ctgatggcac | cgattgtaca | agtataggaa | caatagagca | gaagtctagc | gaagaccaag | 1380 |
| ggataaaggg | tagaattgag | aaagctgcaa | atccaagtgg | caagaagaaa | ctcaagatct | 1440 |
| tccagcctgt | gatagaggcg | cctggtgcct | caaaatgtat | tggccccggg | tgctgtcacg | 1500 |
| tggcgcagcc | cgactcggtg | tactgcagta | atgactgtat | cctcaaacac | gccgcagcga | 1560 |
| caatgaagtt | tctaagctca | ggtaaagaac | agaagccaaa | gcctaaagaa | aagatgaaga | 1620 |
| tgaagccaga | gaagcccagt | cttccgaaat | gcggtgctca | ggcaggtatt | aaaatctctt | 1680 |
| ctgtgcacaa | gagaccagct | ccagaaaaaa | aagagaccac | agtgaagaag | gcagtggtgg | 1740 |
| tccctgcgcg | gagtgaagca | ctcgggaagg | aagcagcttg | tgagagcagc | acgccgtcgt | 1800 |
| gggcgagcga | tcacaattac | aatgcagtaa | agccagaaaa | gactgctgct | ccctcgccgt | 1860 |
| cactgttgta | taaatgtatg | tatcacctag | gggttggcct | cctggacccc | tcccgttctt | 1920 |
| tctggatagc | catcccctgg | gcctgtccag | gactgggagt | tgcagctttg | tgttaagctg | 1980 |
| atcacagaca | ccggctgcac | catcagcggg | aagcagagcc | catgtccagg | atgcctcctg | 2040 |
| ctgccctgtg | tccatcccta | gtctgtcagg | acttcctgtc | actgttttcc | aaagctgtaa | 2100 |
| acctcactgg | tgaacgttca | ccttaatgat | tgattcttta | atctctgttt | tcactctcag | 2160 |
| gctctggtaa | gtattcgtat | tctcttcatc | ccagtctgat | tgcatagcca | cactgcccgg | 2220 |
| cacgccacat | ccacccctgt | ctgcacatga | gttgttctga | caacagcgct | gtatacgctt | 2280 |
| cagtttttcc | acattgtcca | cggccagcac | atgaaagcat | cacttcttt | ttatgttgtg | 2340 |
| ggaatctttg | caagttagtg | ttgcatctga | ttttcaggtg | tacatttatt | tttgactggg | 2400 |
| cagatagggg | attttttt | ttccatgtcc | gattcacacg | ctacacaccc | acatgaacac | 2460 |
| attcgaactt | : cgaaggccac | acactcctgc | ttcataggcc | ccacggtaag | tgagttcaca | 2520 |
| cctagaacac | tgtcctgacc | gcaggacgcg | tgccttggac | ttggtattct | acatgtgact | 2580 |
| ggctttcttg | ccctcgtctc | ttgaatgttt | agactcttaa | gatcatatco | tgccccaaat | 2640 |
| ttcaaattaa | tgaaatgaag | atatttcaaa | cagatctttg | aaacctcaga | ttctgtggtg | 2700 |
| caattttaat | gttttcttgt | : ttctcagttt | : tctgctataa | aactattttc | aattcagtc | 2759 |
| | | | | | | |

<210> 122

<211> 781

<212> DNA

<213> NM_177953.1| Homo sapiens dynein, cytoplasmic, light polypeptide 2A (DNCL2A), transcript variant 2, mRNA $\,$

| <400> 122 cgcagaaagg | cacaggactc | gctaagtgtt | cgctacgcgg | ggctaccgga | tcggtcggaa | 60 |
|----------------------|------------|------------|------------|------------|------------|-----|
| atggctgaag 1 | tggagagatc | gcctgagccc | aggaggtcaa | ggctacagtg | agccgtgact | 120 |
| gcaccactgc a | actccaccct | gggcagaggt | ggaggagaca | ctgaagcgac | tgcagagcca | 180 |
| gaagggagtg | cagggaatca | tcgtcgtgaa | cacagaaggc | attcccatca | agagcaccat | 240 |
| ggacaacccc | | | | | | 300 |
| gagcaccgtg | | | | | | 360 |
| gaaaaatgaa | | | | | | 420 |
| | gccactctct | | | | | 480 |
| | caatcatgtc | | | | | 540 |
| | gaccgtgtgt | | | | | 600 |
| | ttccccagag | | | | | 660 |
| | gaagcccgca | | | | | 720 |
| | gtttttcttt | | | | | 780 |
| | <i>9</i> | | _ | | | 781 |
| a | | | | | | |

<210> 123

<211> 841

<212> DNA

 $<\!\!213\!\!>\!$ NM_022873.1| Homo sapiens interferon, alpha-inducible protein (clone IFI-6-16) (G1P3), transcript variant 3, mRNA

| <400> 123 gaaccgttta ctcgctgctg tgcccatcta tcagcaggct ccgggctgaa ga | ttgcttct 60 |
|--|--------------|
| cttctctcct ccaaggtcta gtgacggagc ccgcgcggg cgccaccatg cg | gcagaagg 120 |
| cggtatcgct tttcttgtgc tacctgctgc tcttcacttg cagtggggtg ga | |
| agaatgcggg taaggatgca ggtaagaaaa agtgctcgga gagctcggac ag | |
| ggttctggaa ggccctgacc ttcatggccg tcggaggagg actcgcagtc gc | |
| ccgcgctggg cttcaccggc gccggcatcg cggccaactc ggtggctgcc tc | |
| gctggtctgc gatcctgaat gggggcggcg tgcccgccgg ggggctagtg gc | |
| agagcctcgg ggctggtggc agcagcgtcg tcataggtaa tattggtgcc ct | |
| acgccacca caagtatctc gatagtgagg aggatgagga gtagccagca gc | - 40 |
| | |

| cctcttcttc | cttcttggcc | taactcttcc | agttaggatc | tagaactttg | ccttttttt | 600 |
|------------|------------|------------|------------|------------|------------|-----|
| tttttttt | ttttttgag | atgggttctc | actatattgt | ccaggctaga | gtgcagtggc | 660 |
| tattcacaga | tgcgaacata | gtacactgca | gcctccaact | cctagcctca | agtgatcctc | 720 |
| ctgtctcaac | ctcccaagta | ggattacaag | catgcgccga | cgatgcccag | aatccagaac | 780 |
| tttgtctatc | actctcccca | acaacctaga | tgtgaaaaca | gaataaactt | cacccagaaa | 840 |
| a | | | | | | 841 |

<210> 124

<211> 4652

<212> DNA

 $<\!\!213\!\!>\!$ NM_183047.1| Homo sapiens protein kinase C binding protein 1 (PRKCBP1), transcript variant 1, mRNA

| <400> 124 gtgagaacta | ggagcctgtc | ctccatgttt | tataagtatt | gacattacac | agtgttaaca | 60 |
|-------------------------|--------------|------------|--------------|------------|------------|------|
| | agagcttggc | | | | | 120 |
| atggatatct | ctactcgctc | caaagatcct | ggctctgcag | agagaacagc | ccagaaaaga | 180 |
| aagttcccca | gccctccaca | ttcttccaat | ggccactcgc | cgcaggacac | atcaacaagc | 240 |
| cccattaaaa | agaaaaagaa | acctggctta | ctgaacagta | acaataagga | gcagtcagaa | 300 |
| ctaagacatg | gtccgtttta | ctatatgaag | cagccactca | ccacagaccc | tgttgatgtt | 360 |
| gtaccgcagg | atggacggaa | tgatttctac | tgctgggttt | gtcaccggga | aggccaagtc | 420 |
| ctttgctgtg | agctctgtcc | ccgggtttat | cacgctaagt | gtctgagact | gacatcggaa | 480 |
| ccagaggggg | actggttttg | tcctgaatgt | gagaaaatta | cagtagcaga | atgcatcgag | 540 |
| acccagagta | aagccatgac | aatgctcacc | attgaacagt | tatcctacct | gctcaagttt | 600 |
| gccattcaga | aaatgaaaca | gccagggaca | gatgcattcc | agaagcccgt | tccattggaa | 660 |
| | actatgcgga | | | | | 720 |
| | agaaaatgta | | | | | 780 |
| ttgcacaact | gcatcattta | taatggggga | aatcacaaat | tgacgcaaat | agcgaaagta | 840 |
| gtcatcaaaa | tctgtgaaca | tgagatgaat | gaaatcgaag | tatgtccaga | atgttatcta | 900 |
| | aaaaacgaga | | | | | 960 |
| gtctgggcca | aactgaaggg | gtttccattc | tggcctgcaa | aagctctaag | ggataaagac | 1020 |
| gggcaggtcg | atgcccgatt | ctttggacaa | catgacaggg | cctgggttcc | aataaataat | 1080 |
| tgctacctca | a tgtctaaaga | aattcctttt | : tctgtgaaaa | agactaagag | catcttcaac | 1140 |
| agtgccatgo | aagagatgga | ggtttacgtg | gagaacatcc | gcaggaagtt | tggggtttt | 1200 |
| aattactct | catttaggac | accctacaca | cccaacagcc | agtatcaaat | gctgctcgat | 1260 |
| | | | | | | |

| cccaccaacc | ccagcgccgg | cactgc c aag | atagacaagc | aggagaaggt | caagctcaac | 1320 |
|------------|------------|---------------------|------------|------------|------------|------|
| tttgacatga | cggcatcccc | caagatcctg | atgagcaagc | ctgtgctgag | tgggggcaca | 1380 |
| ggccgccgga | tttccttgtc | ggatatgccg | cgctccccca | tgagcacaaa | ctcttctgtg | 1440 |
| cacacgggct | ccgacgtgga | gcaggatgct | gagaagaagg | ccacgtcgag | ccacttcagt | 1500 |
| gcgagcgagg | agtccatgga | cttcctggat | aagagcacag | cttcaccagc | ctccaccaag | 1560 |
| acgggacaag | cagggagttt | atccgg c agc | ccaaagccct | tctctcctca | actgtcagct | 1620 |
| cctatcacga | cgaaaacgga | caaaac c tcc | accaccggca | gcatcctgaa | tcttaacctg | 1680 |
| gatcgaagca | aagctgagat | ggatttgaag | gagctgagcg | agtcggtcca | gcaacagtcc | 1740 |
| acccctgttc | ctctcatctc | tcccaagcgc | cagattcgta | gcaggttcca | gctgaatctt | 1800 |
| gacaagacca | tagagagttg | caaagcacaa | ttaggcataa | atgaaatctc | ggaagatgtc | 1860 |
| tatacggccg | tagagcacag | cgattcggag | gattctgaga | agtcagatag | tagcgatagt | 1920 |
| gagtatatca | gtgatgatga | gcagaa g tct | aagaacgagc | cagaagacac | agaggacaaa | 1980 |
| gaaggttgtc | agatggacaa | agagccatct | gctgttaaaa | aaaagcccaa | gcctacaaac | 2040 |
| ccagtggaga | ttaaagagga | gctgaaaagc | acgtcaccag | ccagcgagaa | ggcagaccct | 2100 |
| ggagcagtca | aggacaaggc | cagccctgag | cctgagaagg | acttttccga | aaaggcaaaa | 2160 |
| ccttcacctc | accccataaa | ggataaactg | aagggaaaag | atgagacgga | ttccccaaca | 2220 |
| gtccatttgg | gcctggactc | tgattcagag | agcgaacttg | tcatagattt | aggagaagac | 2280 |
| cattctgggc | gggagggtcg | aaaaaataag | aaggaaccca | aagaaccatc | tcccaaacag | 2340 |
| gatgttgtag | gtaaaactcc | accatc cacg | acggtgggca | gccattctcc | cccggaaaca | 2400 |
| ccggtgctca | cccgctcttc | cgcccaaact | tccgcggctg | gcgccacagc | caccaccagc | 2460 |
| acgtcctcca | cggtcaccgt | cacggccccg | gccccgccg | ccacaggaag | cccagtgaaa | 2520 |
| aagcagaggc | cgcttttacc | gaaggagact | gccccggccg | tgcagcgggt | cgtgtggaac | 2580 |
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| tcctcagtca | acgctgacct | gcccatcgcc | actgcctcag | ctgatgtcgc | cgctgatatt | 2760 |
| gccaagtaca | ctagcaaaat | gatggatgca | ataaaaggaa | caatgacaga | aatatacaac | 2820 |
| gatctttcta | aaaacactac | tggaagcaca | atagctgaga | ttcgcaggct | gaggatcgag | 2880 |
| atagagaagc | tccagtggct | gcaccagcaa | gagctctccg | aaatgaaaca | caacttagag | 2940 |
| ctgaccatgg | cggagatgcg | gcagagcctg | gagcaggagc | gggaccggct | catcgccgag | 3000 |
| gtgaagaagc | agctggagtt | ggagaagcag | caggcggtgg | atgagaccaa | gaagaagcag | 3060 |
| tggtgcgcca | actgcaagaa | ggaggccatc | ttttactgct | gttggaacac | tagctactgt | 3120 |
| gactacccct | gccagcaagc | ccactggcct | gagcacatga | agtcctgcac | ccagtcagct | 3180 |
| actgctcctc | agcaggaagc | ggatg⊂tgag | gtgaacacag | aaacactaaa | taagtcctcc | 3240 |
| caggggagct | cctcgagcac | acaat⊂agca | ccttcagaaa | cggccagcgc | ctccaaagag | 3300 |
| | | | | | | |

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aaggagacgt cagctgagaa aagcaaggag agtggctcga cccttgacct ttctggctcc
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                                                                  3420
gacaagcaac ctgcctatgc cccaaccacc acagaccacc agccgcaccc caactacccc
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qcccagaagt accattcccg gagtaataaa tccagttgga gcagcagtga tgagaagagg
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ggatcgacac gttccgatca caacaccagt accagcacga agagcctcct cccgaaagag
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tctcggctgg acaccttctg ggactagcag tgaatcggga cacaaaccac ccaccccatt
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cttttggtgt aaataatgta caatttgtgg atgtcattga atctagagga ctttcccctt
                                                                  3900
3960
aaaaaaaaaa aaaaagcaac caaccccaac aacaaaaaag aatgttttgg tattggagaa
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                                                                  4080
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                                                                  4260
cacagggaga agctagtttc tttcatgtga ttgaaatgat gactctactc ctaaaaggga
                                                                  4320
aaaaacaata tccttgttta cagaagagaa acaaacaagc cccactcagc tcagtcacag
                                                                  4380
gagagaacac agaaagtctt aggatcatga actctgaaaa aaagagaaac cttatctttg
                                                                  4440
ctttgtggtt cctttaaaca cactcacaca cacttggtca gagatgctgt gcttcttgga
                                                                  4500
agcaaggact caaaggcaag gtgcacgcag aggacgtttg agtctgggat gaagcatgta
                                                                  4560
                                                                  4620
cgtattattt atatgatgga atttcacgtt tttatgtaag catgaaacac aggcagtatg
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agagaaagca aggcccgtca tgctgtccgt ac
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<210> 125

<211> 3217

<212> DNA

<213> NM_017452.1| Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript variant T2, mRNA

| <400> 125 acttcctgcc | gggctgcggg | cgcctgagcg | ctcttcagcg | tttgcgcggc | ggctgcgcgt | 60 |
|----------------------|------------|------------|------------|------------|------------|-----|
| ctctctcggc | tcccgcttcc | tttgaccgcc | tccccccc | ggcccggcgg | cgcccgcctc | 120 |
| ctccacggcc | actccgcctc | ttccctccct | tcgtcccttc | ttcctctccc | ttttttcctt | 180 |
| cttccttccc | ctcctcgccg | ccaccgccca | ggaccgccgg | ccgggggacg | agtccggagc | 240 |

| agcagccaga | gtttattaac | cacttaacct | ctcagaactg | aacaaagaca | acattgttcc | 300 |
|------------|------------|------------|------------|------------|------------|------|
| tggaacgccc | tctttttaaa | aaagaaagca | taacccctac | tgtagaacta | aatgcactgt | 360 |
| gcatgaaact | tggaaaaaaa | ccaatgtata | agcctgttga | cccttactct | cggatgcagt | 420 |
| ccacctataa | ctacaacatg | agaggaggtg | cttatccccc | gaggtacttt | tacccatttc | 480 |
| cagttccacc | tttactttat | caagtggaac | tttctgtggg | aggacagcaa | tttaatggca | 540 |
| aaggaaagac | aagacaggct | gcgaaacacg | atgctgctgc | caaagcgttg | aggatcctgc | 600 |
| agaatgagcc | cctgccagag | aggctggagg | tgaatggaag | agaatccgaa | gaagaaaatc | 660 |
| tcaataaatc | tgaaataagt | caagtgtttg | agattgcact | taaacggaac | ttgcctgtga | 720 |
| atttcgaggt | ggcccgggag | agtggcccac | cccacatgaa | gaactttgtg | accaaggttt | 780 |
| cggttgggga | gtttgtgggg | gaaggtgaag | ggaaaagcaa | gaagatttca | aagaaaaatg | 840 |
| ccgccatagc | tgttcttgag | gagctgaaga | agttaccgcc | cctgcctgca | gttgaacgag | 900 |
| taaagcctag | aatcaaaaag | aaaacaaaac | ccatagtcaa | gccacagaca | agcccagaat | 960 |
| atggccaggg | gatcaatccg | attagccgac | tggcccagat | ccagcaggca | aaaaaggaga | 1020 |
| aggagccaga | gtacacgctc | ctcacagagc | gaggcctccc | gcgccgcagg | gagtttgtga | 1080 |
| tgcaggtgaa | ggttggaaac | cacactgcag | aaggaacggg | caccaacaag | aaggtggcca | 1140 |
| agcgcaatgc | agccgagaac | atgctggaga | tccttggttt | caaagtcccg | cagcggcagc | 1200 |
| ccaccaaacc | cgcactcaag | tcagaggaga | agacacccat | aaagaaacca | ggggatggaa | 1260 |
| gaaaagtaac | cttttttgaa | cctggctctg | gggatgaaaa | tgggactagt | aataaagagg | 1320 |
| atgagttcag | gatgccttat | ctaagtcatc | agcagctgcc | tgctggaatt | cttcccatgg | 1380 |
| tgcccgaggt | cgcccaggct | gtaggagtta | gtcaaggaca | tcacaccaaa | gattttacca | 1440 |
| gggcagctcc | gaatcctgcc | aaggccacgg | taactgccat | gatagcccga | gagttgttgt | 1500 |
| atgggggcac | ctcgcccaca | gccgagacca | ttttaaagaa | taacatctct | tcaggccacg | 1560 |
| taccccatgg | acctctcacg | agaccctctg | agcaactgga | ctatctttcc | agagtccagg | 1620 |
| gattccaggt | tgaatacaaa | gacttcccca | aaaacaacaa | gaacgaattt | gtatctctta | 1680 |
| tcaattgctc | ctctcagcca | cctctgatca | gccatggtat | cggcaaggat | gtggagtcct | 1740 |
| gccatgatat | ggctgcgctg | aacatcttaa | agttgctgtc | tgagttggac | caacaaagta | 1800 |
| cagagatgcc | aagaacagga | aacggaccaa | tgtctgtgtg | tgggaggtgc | tgaacctttt | 1860 |
| ctggccatga | accattataa | aatcccaaca | tatatactga | aaatactgaa | actgctttga | 1920 |
| aaatttggaa | tttctgatac | ctccagtggg | ccgagagaca | cggtgggtaa | aggatgtggg | 1980 |
| cagcagcagg | gaagacaaca | gaaacacaag | gaggcggctg | tggccggctg | gactgtgctg | 2040 |
| gggtttgttg | tgatggccac | tcggtgacct | ggcggtccct | acgcaatagc | agctgcctgt | 2100 |
| ggggaagaag | ggctgcccag | ccagctggtt | ctcccgggac | accagcagat | ccacaccctg | 2160 |
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| tctcaagctg | gctcactcag | acacattggg | acaaaccctg | gacagccatg | ccagagagag | 2280 |
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| | | | | tgcagtaccc | | 2400 |
| cactgctttc | tcttccaaca | gtgatctgta | ttcttagttt | cattattttc | ttttgattga | 2460 |
| tatgacacta | tataaaattt | tcatttgaga | atttctcaat | tgtatctagt | taaatagcac | 2520 |
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| aaattgtgtg | ccctctggtt | cagctgaaac | agtcctggac | tttcaaaaac | cttgaataag | 2700 |
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| ttccattttt | atttcatttt | tatttttgtt | aatgcaaaca | ggacttaaat | gaactttgat | 2820 |
| ctctgtttta | aagattatta | aaaaacattg | tgtatctata | catatggctc | ttgaggactt | 2880 |
| agctttcact | acactacagg | atatgatctc | catgtagtcc | atataaacct | gcagagtgat | 2940 |
| tttccagagt | gctcgatact | gttaattaca | tctccattag | ggctgaaaag | aatgacctac | 3000 |
| gtttctgtat | acagctgtgt | tgcttttgat | gttgtgttac | tgtacacaga | agtgtgtgca | 3060 |
| ctgaggctct | gcgtgtggtc | cgtatggaaa | acctggtagc | cctgcgagtt | aagtactgct | 3120 |
| tccattcatt | gtttacgctg | gaatttttct | ccccatggaa | tgtaagtaaa | acttaagtgt | 3180 |
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<210> 126

<211> 3506

<212> DNA

 $<\!\!213\!\!>$ NM_017453.1| Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript variant T3, mRNA

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| ctccacggcc | actccgcctc | ttccctccct | tcgtcccttc | ttcctctccc | tttttcctt | 180 |
| cttccttccc | ctcctcgccg | ccaccgccca | ggaccgccgg | ccgggggacg | agtccggagc | 240 |
| agcagccaga | gtttattaac | cacttaacct | ctcagaactg | aacaaagaca | acattgttcc | 300 |
| tggaacgccc | tctttttaaa | aaaggtagaa | ctttagactt | catagcactg | aattaacctg | 360 |
| cactgaaagc | tgtttacctg | catttgttca | cttttgttga | aagtgaccat | gtctcaagtt | 420 |
| caagtgcaag | ttcagaaccc | atctgctgct | ctctcaggga | gccaaatact | gaacaagaac | 480 |
| cagtctcttc | tctcacagcc | tttgatgagt | attccttcta | ctactagctc | tctgccctct | 540 |
| gaaaatgcag | gtagacccat | tcaaaactct | gctttaccct | ctgcatctat | tacatccacc | 600 |

| agtgcagctg | cagaaagcat | aacccctact | gtagaactaa | atgcactgtg | catgaaactt | 660 |
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| tacaacatga | gaggaggtgc | ttatcccccg | aggtactttt | acccatttcc | agttccacct | 780 |
| ttactttatc | aagtggaact | ttctgtggga | ggacagcaat | ttaatggcaa | aggaaagaca | 840 |
| agacaggctg | cgaaacacga | tgctgctgcc | aaagcgttga | ggatcctgca | gaatgagccc | 900 |
| ctgccagaga | ggctggaggt | gaatggaaga | gaatccgaag | aagaaaatct | caataaatct | 960 |
| gaaataagtc | aagtgtttga | gattgcactt | aaacggaact | tgcctgtgaa | tttcgaggtg | 1020 |
| gcccgggaga | gtggcccacc | ccacatgaag | aactttgtga | ccaaggtttc | ggttggggag | 1080 |
| tttgtggggg | aaggtgaagg | gaaaagcaag | aagatttcaa | agaaaaatgc | cgccatagct | 1140 |
| gttcttgagg | agctgaagaa | gttaccgccc | ctgcctgcag | ttgaacgagt | aaagcctaga | 1200 |
| atcaaaaaga | aaacaaaacc | catagtcaag | ccacagacaa | gcccagaata | tggccagggg | 1260 |
| atcaatccga | ttagccgact | ggcccagatc | cagcaggcaa | aaaaggagaa | ggagccagag | 1320 |
| tacacgctcc | tcacagagcg | aggcctcccg | cgccgcaggg | agtttgtgat | gcaggtgaag | 1380 |
| gttggaaacc | acactgcaga | aggaacgggc | accaacaaga | aggtggccaa | gcgcaatgca | 1440 |
| gccgagaaca | tgctggagat | ccttggtttc | aaagtcccgc | agcggcagcc | caccaaaccc | 1500 |
| gcactcaagt | cagaggagaa | gacacccata | aagaaaccag | gggatggaag | aaaagtaacc | 1560 |
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| aatcctgcca | aggccacggt | aactgccatg | atagcccgag | agttgttgta | tgggggcacc | 1800 |
| tcgcccacag | ccgagaccat | tttaaagaat | aacatctctt | caggccacgt | accccatgga | 1860 |
| cctctcacga | gaccctctga | gcaactggac | tatctttcca | gagtccaggg | attccaggtt | 1920 |
| gaatacaaag | acttccccaa | aaacaacaag | aacgaatttg | tatctcttat | caattgctcc | 1980 |
| tctcagccac | ctctgatcag | ccatggtatc | ggcaaggatg | tggagtcctg | ccatgatatg | 2040 |
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| agaacaggaa | acggaccaat | gtctgtgtgt | gggaggtgct | gaaccttttc | tggccatgaa | 2160 |
| ccattataaa | atcccaacat | atatactgaa | aatactgaaa | ctgctttgaa | aatttggaat | 2220 |
| ttctgatacc | tccagtgggc | cgagagacac | ggtgggtaaa | ggatgtgggc | agcagcaggg | 2280 |
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| gccccagagc | taaaagcacc | agagaaaatc | aaatgcttcc | tactcagcgt | gacccaactt | 2640 |

| ttctagtgtg | ccacggcccc | accacctcct | gcagtaccca | caccatcacc | actgctttct | 2700 |
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| cttccaacag | tgatctgtat | tcttagtttc | attattttct | tttgattgat | atgacactat | 2760 |
| ataaaatttt | catttgagaa | tttctcaatt | gtatctagtt | aaatagcaca | gtttggaaac | 2820 |
| ttgtctgaga | ctgactttat | caataatcta | accgacaaag | atcatatcca | tgtgtatgtg | 2880 |
| gttagacatt | tttatttcat | tgactaaccc | aggacagttt | cagtgatgca | aattgtgtgc | 2940 |
| cctctggttc | agctgaaaca | gtcctggact | ttcaaaaacc | ttgaataagt | ctcccacagt | 3000 |
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| agattattaa | aaaacattgt | gtatctatac | atatggctct | tgaggactta | gctttcacta | 3180 |
| cactacagga | tatgatctcc | atgtagtcca | tataaacctg | cagagtgatt | ttccagagtg | 3240 |
| ctcgatactg | ttaattacat | ctccattagg | gctgaaaaga | atgacctacg | tttctgtata | 3300 |
| cagctgtgtt | gcttttgatg | ttgtgttact | gtacacagaa | gtgtgtgcac | tgaggctctg | 3360 |
| cgtgtggtcc | gtatggaaaa | cctggtagcc | ctgcgagtta | agtactgctt | ccattcattg | 3420 |
| tttacgctgg | aatttttctc | cccatggaat | gtaagtaaaa | cttaagtgtt | tgtcatcaat | 3480 |
| aaatggtaat | actaaaaaaa | aaaaaa | | | | 3506 |
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<210> 127

<211> 4538

<212> DNA

 $<\!\!213\!\!>$ NM_199169.1| Homo sapiens transmembrane, prostate androgen induced RNA (TMEPAI), transcript variant 2, mRNA

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| 120 | cg g agctgga | caggcaatgg | cctgcgaaac | cggacagtct | gaactgaagg | agctgctgga |
| 180 | tgatcacgtg | atggtggtgg | gatgatggtg | tcgtggtggt | atcatcatca | gtttgttcag |
| 240 | gc c aggggcg | agccggcaca | gtccttcatc | tgtctgcacg | cactacaagc | cctgctgagc |
| 300 | gcacagtgtc | ccctcggaga | atgcctgtgg | cctcagaagg | gatgccctgt | gaggagagaa |
| 360 | accgcctggc | cggcccaccg | cgccccgcct | cgcaggtcta | atcccagagc | aggcaacgga |
| 420 | at ccgtacct | cagcccacct | ccaccgcttc | gggagcgctt | ttcgcccagc | cgtgccgccc |
| 480 | ag cccccacc | gacggggagg | ctcgctgtca | cacccaccat | atcgacctgc | gcagcacgag |
| 540 | aactgaaccg | cagcagctgg | ggaccccgag | tccagcttcg | ccctgcaccc | ctaccagggc |
| 600 | tggatagtgc | agtgacctga | catcttcgac | caaacagaac | cgcgcacccc | ggagtcggtg |
| 660 | cgtgctacgg | atcagcgcca | taactcgggc | ccccagcag | ggcccctgcc | caggctgggc |
| 720 | gccactaccc | gaggtcatcg | cacctacagc | ggccgccgcc | cgcatggagg | cagcggcggg |
| | | | | | | |

| ggggtcctcc | ttccagcacc | agcagagcag | tgggccgccc | tccttgctgg | aggggacccg | 780 |
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| gctccaccac | acacacatcg | cgcccctaga | gagcgcagcc | atctggagca | aagagaagga | 840 |
| taaacagaaa | ggacaccctc | tctagggtcc | ccaggggggc | cgggctgggg | ctgcgtaggt | 900 |
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| aaccacgttt | ctttgttgag | ctgtgtcttg | aaggcaaaag | aaaaaaaatt | tctacagtag | 1140 |
| tctttcttgt | ttctagttga | gctgcgtgcg | tgaatgctta | ttttcttttg | tttatgataa | 1200 |
| tttcacttaa | ctttaaagac | atatttgcac | aaaacctttg | tttaaagatc | tgcaatatta | 1260 |
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| gagttccagg | tagtggtttt | gcctttccca | aaaatgaaaa | taaactgtta | ccgaaggaat | 1860 |
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| ctcacaagcc | gcagcggctg | gtgcccgggc | taccagggac | atgccagagg | gctcgatgac | 1980 |
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| aatgctttca | cgttggttcc | tggcgtggga | actgctctcc | tttgcagccc | catttcccaa | 4320 |
| gctctgttca | agttaaactt | atgtaagctt | tccgtggcat | gcggggcgcg | cacccacgtc | 4380 |
| cccgctgcgt | aagactctgt | atttggatgc | caatccacag | gcctgaagaa | actg⊂ttgtt | 4440 |
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<210> 128

<211> 4531

<212> DNA

 $<\!\!213\!\!>\,$ NM_199170.1| Homo sapiens transmembrane, prostate androgen induced RNA (TMEPAI), transcript variant 3, mRNA

| <400> 128 tggtcgtcct | ccttgggttc | gggtgaaagc | gcttgggggt | tcagtgggcc | atgatccccg | 60 |
|----------------------|------------|------------|------------|------------|------------|------|
| agctgctgga | gaactgaagg | cggacagtct | cctgcgaaac | cagcggagct | ggagtttgtt | 120 |
| cagatcatca | tcatcgtggt | ggtgatgatg | gtgatggtgg | tggtgatcac | gtgcctgctg | 180 |
| agccactaca | agctgtctgc | acggtccttc | atcagccggc | acagccaggg | gcggaggaga | 240 |
| gaagatgccc | tgtcctcaga | aggatgcctg | tggccctcgg | agagcacagt | gtcaggcaac | 300 |
| ggaatcccag | agccgcaggt | ctacgccccg | cctcggccca | ccgaccgcct | ggccgtgccg | 360 |
| cccttcgccc | agcgggagcg | cttccaccgc | ttccagccca | cctatccgta | cctgcagcac | 420 |
| gagatcgacc | tgccacccac | catctcgctg | tcagacgggg | aggagccccc | accctaccag | 480 |
| ggcccctgca | ccctccagct | tcgggacccc | gagcagcagc | tggaactgaa | ccgggagtcg | 540 |
| gtgcgcgcac | ccccaaacag | aaccatcttc | gacagtgacc | tgatggatag | tgccaggctg | 600 |
| ggcggcccct | gccccccag | cagtaactcg | ggcatcagcg | ccacgtgcta | cggcagcggc | 660 |
| gggcgcatgg | aggggccgcc | gcccacctac | agcgaggtca | tcggccacta | cccggggtcc | 720 |
| tccttccagc | accagcagag | cagtgggccg | ccctccttgc | tggaggggac | ccggctccac | 780 |
| cacacacaca | tcgcgcccct | agagagcgca | gccatctgga | gcaaagagaa | ggataaacag | 840 |
| aaaggacacc | ctctctaggg | tccccagggg | ggccgggctg | gggctgcgta | ggtgaaaagg | 900 |
| cagaacactc | cgcgcttctt | agaagaggag | tgagaggaag | gcggggggcg | cagcaacgca | 960 |
| tcgtgtggcc | ctccctccc | acctccctgt | gtataaatat | ttacatgtga | tgtctggtct | 1020 |
| gaatgcacaa | gctaagagag | cttgcaaaaa | aaaaaagaaa | aaagaaaaaa | aaaaaccacg | 1080 |
| tttctttgtt | gagctgtgtc | ttgaaggcaa | aagaaaaaaa | atttctacag | tagtctttct | 1140 |
| tgtttctagt | tgagctgcgt | gcgtgaatgc | ttattttctt | ttgtttatga | taatttcact | 1200 |
| taactttaaa | gacatatttg | cacaaaacct | ttgtttaaag | atctgcaata | ttatatatat | 1260 |
| aaatatatat | aagataagag | aaactgtatg | tgcgagggca | ggagtatttt | tgtattagaa | 1320 |
| gaggcctatt | aaaaaaaaa | gttgttttct | gaactagaag | aggaaaaaaa | tggcaatttt | 1380 |
| tgagtgccaa | gtcagaaagt | gtgtattacc | ttgtaaagaa | aaaaattaca | aagcaggggt | 1440 |
| ttagagttat | ttatataaat | gttgagattt | tgcactattt | tttaatataa | atatgtcagt | 1500 |
| gcttgcttga | tggaaacttc | tcttgtgtct | gttgagactt | taagggagaa | atgtcggaat | 1560 |
| ttcagagtcg | cctgacggca | gagggtgagc | ccccgtggag | tctgcagaga | ggccttggcc | 1620 |
| aggagcggcg | ggctttcccg | aggggccact | gtccctgcag | agtggatgct | tctgcctagt | 1680 |
| gacaggttat | caccacgtta | tatattccct | accgaaggag | acaccttttc | cccctgacc | 1740 |
| cagaacagco | tttaaatcac | aagcaaaata | ggaaagttaa | ccacggaggc | accgagttcc | 1800 |

| aggtagtggt | tttgcctttc | ccaaaaatga | aaataaactg | ttaccgaagg | aattagtttt | 1860 |
|------------|------------|------------|------------|------------|------------|--------------|
| tcctcttctt | ttttccaact | gtgaaggtcc | ccgtggggtg | gagcatggtg | ccctcacaa | 1920 |
| gccgcagcgg | ctggtgcccg | ggctaccagg | gacatgccag | agggctcgat | gacttgtctc | 1980 |
| tgcagggcgc | tttggtggtt | gttcagctgg | ctaaaggttc | accggtgaag | gcaggtgcgg | 2040 |
| taactgccgc | actggaccct | aggaagcccc | aggtattcgc | aatctgacct | cctcctgtct | 2100 |
| gtttcccttc | acggatcaat | tctcacttaa | gaggccaata | aacaacccaa | catgaaaagg | 2160 |
| tgacaagcct | gggtttctcc | caggataggt | gaaagggtta | aaatgagtaa | agcagttgag | 2220 |
| caaacaccaa | cccgagcttc | gggcgcagaa | ttcttcacct | tctcttcccc | tttccatctc | 2280 |
| ctttccccgc | ggaaacaacg | cttcccttct | ggtgtgtctg | ttgatctgtg | ttttcattta | 2340 |
| catctctctt | agactccgct | cttgttctcc | aggttttcac | cagatagatt | tggggttggc | 2400 |
| gggacctgct | ggtgacgtgc | aggtgaagga | caggaagggg | catgtgagcg | taaatagagg | 2460 |
| tgaccagagg | agagcatgag | gggtggggct | ttgggaccca | ccggggccag | tggctggagc | 2520 |
| ttgacgtctt | tcctccccat | gggggtggga | gggcccccag | ctggaagagc | agactcccag | 2580 |
| ctgctacccc | ctcccttccc | atgggagtgg | ctttccattt | tgggcagaat | gctgactagt | 2640 |
| agactaacat | aaaagatata | aaaggcaata | actattgttt | gtgagcaact | tttttataac | 2700 |
| ttccaaaaca | aaaacctgag | cacagttttg | aagttctagc | cactcgagct | catgcatgtg | 2760 |
| aaacgtgtgc | tttacgaagg | tggcagctga | cagacgtggg | ctctgcatgc | cgccagccta | 2820 |
| gtagaaagtt | ctcgttcatt | ggcaacagca | gaacctgcct | ctccgtgaag | tcgtcagcct | 2880 |
| aaaatttgtt | tctctcttga | agaggattct | ttgaaaaggt | cctgcagaga | aatcagtaca | 2940 |
| ggttatcccg | aaaggtacaa | ggacgcactt | gtaaagatga | ttaaaacgta | tctttccttt | 3000 |
| atgtgacgcg | tctctagtgc | cttactgaag | aagcagtgac | actcccgtcg | ctcggtgagg | 3060 |
| acgttcccgg | acagtgcctc | actcacctgg | gactggtatc | ccctcccagg | gtccaccaag | 31.20 |
| ggctcctgct | tttcagacac | cccatcatcc | tcgcgcgtcc | tcaccctgtc | tctaccaggg | 31 80 |
| aggtgcctag | cttggtgagg | ttactcctgc | tcctccaacc | tttttttgcc | aaggtttgta | 3240 |
| cacgactccc | atctaggctg | aaaacctaga | agtggacctt | gtgtgtgtgc | atggtgtcag | 3300 |
| cccaaagcca | ggctgagaca | gtcctcatat | cctcttgagc | caaactgttt | gggtctcgtt | 3360 |
| gcttcatggt | atggtctgga | tttgtgggaa | tggctttgcg | tgagaaaggg | gaggagagtg | 3420 |
| gttgctgccc | tcagccggct | tgaggacaga | gcctgtccct | ctcatgacaa | ctcagtgttg | 3480 |
| aagcccagtg | tcctcagctt | catgtccagt | ggatggcaga | agttcatggg | gtagtggcct | 3540 |
| ctcaaaggct | gggcgcatcc | caagacagcc | agcaggttgt | ctctggaaac | gaccagagtt | 3600 |
| aagctctcgg | cttctctgct | gagggtgcac | cctttcctct | agatggtagt | tgtcacgtta | 3660 |
| tctttgaaaa | ctcttggact | gctcctgagg | aggccctctt | ttccagtagg | aagttagatg | 3720 |
| ggggttctca | gaagtggctg | attggaaggg | gacaagcttc | gtttcagggg | tctgccgttc | 3780 |
| | | | | | | |

| catcctggtt | cagagaaggc | cgagcgtggc | tttctctagc | cttgtcactg | tctccctgcc | 3840 |
|------------|------------|------------|------------|------------|------------|------|
| tgtcaatcac | cacctttcct | ccagaggagg | aaaattatct | cccctgcaaa | gcccggttct | 3900 |
| acacagattt | cacaaattgt | gctaagaacc | gtccgtgttc | tcagaaagcc | cagtgttttt | 3960 |
| gcaaagaatg | aaaagggacc | ccatatgtag | caaaaatcag | ggctggggga | gagccgggtt | 4020 |
| cattccctgt | cctcattggt | cgtccctatg | aattgtacgt | ttcagagaaa | tttttttcc | 4080 |
| tatgtgcaac | acgaagcttc | cagaaccata | aaatatcccg | tcgataagga | aagaaaatgt | 4140 |
| cgttgttgtt | gtttttctgg | aaactgcttg | aaatcttgct | gtactataga | gctcagaagg | 4200 |
| acacagcccg | tcctcccctg | cctgcctgat | tccatggctg | ttgtgctgat | tccaatgctt | 4260 |
| tcacgttggt | tcctggcgtg | ggaactgctc | tcctttgcag | ccccatttcc | caagctctgt | 4320 |
| tcaagttaaa | cttatgtaag | ctttccgtgg | catgcggggc | gcgcacccac | gtccccgctg | 4380 |
| cgtaagactc | tgtatttgga | tgccaatcca | caggcctgaa | gaaactgctt | gttgtgtatc | 4440 |
| agtaatcatt | agtggcaatg | atgacattct | gaaaagctgc | aatacttata | caataaattt | 4500 |
| tacaattctt | tggaaaaaaa | aaaaaaaaa | a | | | 4531 |
| | | | | | | |

<210> 129

<211> 2692

<212> DNA

 $<\!\!213\!\!>$ NM_152871.1| Homo sapiens tumor necrosis factor receptor superfamily, member 6 (TNFRSF6), transcript variant 2, mRNA

| . 60 | cgggcgttcc | ctccccaacc | caatggagcc | gttgctgaat | cgcaggccaa | <400> 129 cctacccgcg |
|------|------------|------------|------------|------------|------------|----------------------|
| 120 | tcgtctctga | tttcgtgagc | caccggggct | tcctcctgac | ttccttccca | ccagcgaggc |
| 180 | gggaagcggt | tggggagtga | aagacgcttc | caggtgttca | gagtgacaca | tctcgcgcaa |
| 240 | accctgaggc | cacggaacac | cgggcactgg | gcctcagggg | cttggctgga | ttacgagtga |
| 300 | cccgctcagt | ggttggtgga | ttctcccgcg | gagctgcctc | tgcccaggcg | cagccctggc |
| 360 | tgggcatctg | acaaccatgc | ggattgctca | tcacttcgga | ggaagctctt | acggagttgg |
| 420 | gtgttaatgc | tcgtccaaaa | tgctagatta | ttacgtctgt | cctctggttc | gaccctccta |
| 480 | ctacagttga | aagactgtta | ggaattgagg | ccaagggatt | gacatcaact | ccaagtgact |
| 540 | cctgtcctcc | tgccataagc | tggccaattc | tgcatcatga | ttggaaggcc | gactcagaac |
| 600 | gcgtgccctg | gaaccagact | caatggggat | actgcacagt | aaagctaggg | aggtgaaagg |
| 660 | gaagatgtag | tccaaatgca | ccatttttct | cagacaaagc | aaggagtaca | ccaagaaggg |
| 720 | cccagaatac | tgcacccgga | ggaaataaac | gcttagaagt | gaaggacatg | attgtgtgat |
| 780 | actgtgaccc | gtatgtgaac | taactctact | actttttttg | tgtaaaccaa | caagtgcaga |
| 840 | acaccaagtg | ctcaccagca | ggaatgcaca | gaatcatcaa | tgtgaacatg | ttgcaccaaa |

| caaagaggaa | gtgaagagaa | aggaagtaca | gaaaacatgc | agaaagcaca | gaaaggaaaa | 900 |
|------------|------------|------------|------------|------------|------------|------|
| ccaaggttct | catgaatctc | caaccttaaa | tcctgaaaca | gtggcaataa | atttatctga | 960 |
| tgttgacttg | agtaaatata | tcaccactat | tgctggagtc | atgacactaa | gtcaagttaa | 1020 |
| aggctttgtt | cgaaagaatg | gtgtcaatga | agccaaaata | gatgagatca | agaatgacaa | 1080 |
| tgtccaagac | acagcagaac | agaaagttca | actgcttcgt | aattggcatc | aacttcatgg | 1140 |
| aaagaaagaa | gcgtatgaca | cattgattaa | agatctcaaa | aaagccaatc | tttgtactct | 1200 |
| tgcagagaaa | attcagacta | tcatcctcaa | ggacattact | agtgactcag | aaaattcaaa | 1260 |
| cttcagaaat | gaaatccaaa | gcttggtcta | gagtgaaaaa | caacaaattc | agttctgagt | 1320 |
| atatgcaatt | agtgtttgaa | aagattctta | atagctggct | gtaaatactg | cttggttttt | 1380 |
| tactgggtac | attttatcat | ttattagcgc | tgaagagcca | acatatttgt | agatttttaa | 1440 |
| tatctcatga | ttctgcctcc | aaggatgttt | aaaatctagt | tgggaaaaca | aacttcatca | 1500 |
| agagtaaatg | cagtggcatg | ctaagtaccc | aaataggagt | gtatgcagag | gatgaaagat | 1560 |
| taagattatg | ctctggcatc | taacatatga | ttctgtagta | tgaatgtaat | cagtgtatgt | 1620 |
| tagtacaaat | gtctatccac | aggctaaccc | cactctatga | atcaatagaa | gaagctatga | 1680 |
| ccttttgctg | aaatatcagt | tactgaacag | gcaggccact | ttgcctctaa | attacctctg | 1740 |
| ataattctag | agattttacc | atatttctaa | actttgttta | taactctgag | aagatcatat | 1800 |
| ttatgtaaag | tatatgtatt | tgagtgcaga | atttaaataa | ggctctacct | caaagacctt | 1860 |
| tgcacagttt | attggtgtca | tattatacaa | tatttcaatt | gtgaattcac | atagaaaaca | 1920 |
| ttaaattata | atgtttgact | attatatatg | tgtatgcatt | ttactggctc | aaaactacct | 1980 |
| acttctttct | caggcatcaa | aagcattttg | agcaggagag | tattactaga | gctttgccac | 2040 |
| ctctccattt | ttgccttggt | gctcatctta | atggcctaat | gcacccccaa | acatggaaat | 2100 |
| atcaccaaaa | aatacttaat | agtccaccaa | aaggcaagac | tgcccttaga | aattctagcc | 2160 |
| tggtttggag | atactaactg | ctctcagaga | aagtagcttt | gtgacatgtc | atgaacccat | 2220 |
| gtttgcaatc | aaagatgata | aaatagattc | ttatttttcc | cccacccccg | aaaatgttca | 2280 |
| ataatgtccc | atgtaaaacc | tgctacaaat | ggcagcttat | acatagcaat | ggtaaaatca | 2340 |
| tcatctggat | ttaggaattg | ctcttgtcat | acccccaagt | ttctaagatt | taagattctc | 2400 |
| cttactacta | tcctacgttt | aaatatcttt | gaaagtttgt | attaaatgtg | aattttaaga | 2460 |
| aataatattt | atatttctgt | aaatgtaaac | tgtgaagata | gttataaact | gaagcagata | 2520 |
| cctggaacca | cctaaagaac | ttccatttat | ggaggatttt | tttgcccctt | gtgtttggaa | 2580 |
| ttataaaata | taggtaaaag | tacgtaatta | aataatgttt | ttggtaaaaa | aaaaaaaaa | 2640 |
| aaaaaaaaa | aaaaaaaaaa | aaaaaaaaa | aaaaaaaaa | aaaaaaaaaa | aa | 2692 |
| | | | | | | |

<210> 130

<211> 2730

<212> DNA

 $<\!\!213\!\!>$ NM_152872.1| Homo sapiens tumor necrosis factor receptor superfamily, member 6 (TNFRSF6), transcript variant 3, mRNA

| <400> 130 cctacccgcg | cgcaggccaa | gttgctgaat | caatggagcc | ctccccaacc | cgggcgttcc | 60 |
|----------------------|------------|------------|------------|------------|------------|------|
| ccagcgaggc | ttccttccca | tcctcctgac | caccggggct | tttcgtgagc | tcgtctctga | 120 |
| tctcgcgcaa | gagtgacaca | caggtgttca | aagacgcttc | tggggagtga | gggaagcggt | 180 |
| ttacgagtga | cttggctgga | gcctcagggg | cgggcactgg | cacggaacac | accctgaggc | 240 |
| cagccctggc | tgcccaggcg | gagctgcctc | ttctcccgcg | ggttggtgga | cccgctcagt | 300 |
| acggagttgg | ggaagctctt | tcacttcgga | ggattgctca | acaaccatgc | tgggcatctg | 360 |
| gaccctccta | cctctggttc | ttacgtctgt | tgctagatta | tcgtccaaaa | gtgttaatgc | 420 |
| ccaagtgact | gacatcaact | ccaagggatt | ggaattgagg | aagactgtta | ctacagttga | 480 |
| gactcagaac | ttggaaggcc | tgcatcatga | tggccaattc | tgccataagc | cctgtcctcc | 540 |
| aggtgaaagg | aaagctaggg | actgcacagt | caatggggat | gaaccagact | gcgtgccctg | 600 |
| ccaagaaggg | aaggagtaca | cagacaaagc | ccatttttct | tccaaatgca | gaagatgtag | 660 |
| attgtgtgat | gaaggacatg | gcttagaagt | ggaaataaac | tgcacccgga | cccagaatac | 720 |
| caagtgcaga | tgtaaaccaa | actttttttg | taactctact | gtatgtgaac | actgtgaccc | 780 |
| ttgcaccaaa | tgtgaacatg | gaatcatcaa | ggaatgcaca | ctcaccagca | acaccaagtg | 840 |
| caaagaggaa | ggatccagat | ctaacttggg | gtggctttgt | cttcttcttt | tgccaattcc | 900 |
| actaattgtt | tgggtgaaga | gaaaggaagt | acagaaaaca | tgcagaaagc | acagaaagga | 960 |
| aaaccaaggt | tctcatgaat | ctccaacctt | aaatcctatg | ttgacttgag | taaatatatc | 1020 |
| accactattg | ctggagtcat | gacactaagt | caagttaaag | gctttgttcg | aaagaatggt | 1080 |
| gtcaatgaag | ccaaaataga | tgagatcaag | aatgacaatg | tccaagacac | agcagaacag | 1140 |
| aaagttcaac | tgcttcgtaa | ttggcatcaa | cttcatggaa | agaaagaagc | gtatgacaca | 1200 |
| ttgattaaag | atctcaaaaa | agccaatctt | tgtactcttg | cagagaaaat | tcagactatc | 1260 |
| atcctcaagg | acattactag | tgactcagaa | aattcaaact | tcagaaatga | aatccaaagc | 1320 |
| ttggtctaga | gtgaaaaaca | acaaattcag | ttctgagtat | atgcaattag | tgtttgaaaa | 1380 |
| gattcttaat | agctggctgt | aaatactgct | tggttttta | ctgggtacat | tttatcattt | 1440 |
| attagcgctg | aagagccaac | atatttgtag | atttttaata | tctcatgatt | ctgcctccaa | 1500 |
| ggatgtttaa | aatctagttg | ggaaaacaaa | cttcatcaag | agtaaatgca | gtggcatgct | 1560 |
| aagtacccaa | ataggagtgt | atgcagagga | tgaaagatta | agattatgct | ctggcatcta | 1620 |
| acatatgatt | ctgtagtatg | aatgtaatca | gtgtatgtta | gtacaaatgt | ctatccacag | 1680 |
| gctaacccca | ctctatgaat | caatagaaga | agctatgacc | ttttgctgaa | atatcagtta | 1740 |
| ctgaacaggc | aggccacttt | gcctctaaat | tacctctgat | aattctagag | attttaccat | 1800 |
| | | | | | | |

| atttctaaac | tttgtttata | actctgagaa | gatcatattt | atgtaaagta | tatgtatttg | 1860 |
|------------|------------|------------|------------|------------|------------|------|
| agtgcagaat | ttaaataagg | ctctacctca | aagacctttg | cacagtttat | tggtgtcata | 1920 |
| ttatacaata | tttcaattgt | gaattcacat | agaaaacatt | aaattataat | gtttgactat | 1980 |
| tatatatgtg | tatgcatttt | actggctcaa | aactacctac | ttctttctca | ggcatcaaaa | 2040 |
| gcattttgag | caggagagta | ttactagagc | tttgccacct | ctccattttt | gccttggtgc | 2100 |
| tcatcttaat | ggcctaatgc | acccccaaac | atggaaatat | caccaaaaaa | tacttaatag | 2160 |
| tccaccaaaa | ggcaagactg | cccttagaaa | ttctagcctg | gtttggagat | actaactgct | 2220 |
| ctcagagaaa | gtagctttgt | gacatgtcat | gaacccatgt | ttgcaatcaa | agatgataaa | 2280 |
| atagattctt | atttttcccc | cacccccgaa | aatgttcaat | aatgtcccat | gtaaaacctg | 2340 |
| ctacaaatgg | cagcttatac | atagcaatgg | taaaatcatc | atctggattt | aggaattgct | 2400 |
| cttgtcatac | ccccaagttt | ctaagattta | agattctcct | tactactatc | ctacgtttaa | 2460 |
| atatctttga | aagtttgtat | taaatgtgaa | ttttaagaaa | taatatttat | atttctgtaa | 2520 |
| atgtaaactg | tgaagatagt | tataaactga | agcagatacc | tggaaccacc | taaagaactt | 2580 |
| ccatttatgg | aggattttt | tgccccttgt | gtttggaatt | ataaaatata | ggtaaaagta | 2640 |
| cgtaattaaa | taatgttttt | ggtaaaaaaa | aaaaaaaaa | aaaaaaaaa | aaaaaaaaa | 2700 |
| aaaaaaaaa | aaaaaaaaa | aaaaaaaaa | | | | 2730 |
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<210> 131

<211> 2563

<212> DNA

 $<\!\!213\!\!>$ NM_152874.1| Homo sapiens tumor necrosis factor receptor superfamily, member 6 (TNFRSF6), transcript variant 8, mRNA

| | | | | | | 100 131 |
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| 60 | cgggcgttcc | ctcccaacc | caatggagcc | gttgctgaat | cgcaggccaa | <400> 131 cctacccgcg |
| 120 | tcgtctctga | tttcgtgagc | caccggggct | tcctcctgac | ttccttccca | ccagcgaggc |
| 180 | gggaagcggt | tggggagtga | aagacgcttc | caggtgttca | gagtgacaca | tctcgcgcaa |
| 240 | accctgaggc | cacggaacac | cgggcactgg | gcctcagggg | cttggctgga | ttacgagtga |
| 300 | cccgctcagt | ggttggtgga | ttctcccgcg | gagctgcctc | tgcccaggcg | cagccctggc |
| 360 | tgggcatctg | acaaccatgc | ggattgctca | tcacttcgga | ggaagctctt | acggagttgg |
| 420 | gtgttaatgc | tcgtccaaaa | tgctagatta | ttacgtctgt | cctctggttc | gaccctccta |
| 480 | ctacagttga | aagactgtta | ggaattgagg | ccaagggatt | gacatcaact | ccaagtgact |
| 540 | cctgtcctcc | tgccataagc | tggccaattc | tgcatcatga | ttggaaggcc | gactcagaac |
| 600 | gcgtgccctg | gaaccagact | caatggggat | actgcacagt | aaagctaggg | aggtgaaagg |
| 660 | gaagatgtag | tccaaatgca | ccatttttct | cagacaaagc | aaggagtaca | ccaagaaggg |

| attgtgtgat | gaaggacatg | atgtgaacat | ggaatcatca | aggaatgcac | actcaccagc | 720 |
|------------|------------|------------|------------|------------|------------|--------------|
| aacaccaagt | gcaaagagga | aggatccaga | tctaacttgg | ggtggctttg | tcttcttctt | 780 |
| ttgccaattc | cactaattgt | ttggggaaac | agtggcaata | aatttatctg | atgttgactt | 840 |
| gagtaaatat | atcaccacta | ttgctggagt | catgacacta | agtcaagtta | aaggctttgt | 900 |
| tcgaaagaat | ggtgtcaatg | aagccaaaat | agatgagatc | aagaatgaca | atgtccaaga | 960 |
| cacagcagaa | cagaaagttc | aactgcttcg | taattggcat | caacttcatg | gaaagaaaga | 1020 |
| agcgtatgac | acattgatta | aagatctcaa | aaaagccaat | ctttgtactc | ttgcagagaa | 1080 |
| aattcagact | atcatcctca | aggacattac | tagtgactca | gaaaattcaa | acttcagaaa | 1 140 |
| tgaaatccaa | agcttggtct | agagtgaaaa | acaacaaatt | cagttctgag | tatatgcaat | 1 200 |
| tagtgtttga | aaagattctt | aatagctggc | tgtaaatact | gcttggtttt | ttactgggta | 1 260 |
| cattttatca | tttattagcg | ctgaagagcc | aacatatttg | tagattttta | atatctcatg | 1 320 |
| attctgcctc | caaggatgtt | taaaatctag | ttgggaaaac | aaacttcatc | aagagtaaat | 1 380 |
| gcagtggcat | gctaagtacc | caaataggag | tgtatgcaga | ggatgaaaga | ttaagattat | 1440 |
| gctctggcat | ctaacatatg | attctgtagt | atgaatgtaa | tcagtgtatg | ttagtacaaa | 1500 |
| tgtctatcca | caggctaacc | ccactctatg | aatcaataga | agaagctatg | accttttgct | 1 560 |
| gaaatatcag | ttactgaaca | ggcaggccac | tttgcctcta | aattacctct | gataattcta | 1620 |
| gagattttac | catatttcta | aactttgttt | ataactctga | gaagatcata | tttatgtaaa | 1 680 |
| gtatatgtat | ttgagtgcag | aatttaaata | aggctctacc | tcaaagacct | ttgcacagtt | 1740 |
| tattggtgtc | atattataca | atatttcaat | tgtgaattca | catagaaaac | attaaattat | 1800 |
| aatgtttgac | tattatatat | gtgtatgcat | tttactggct | caaaactacc | tacttctttc | 1860 |
| tcaggcatca | aaagcatttt | gagcaggaga | gtattactag | agctttgcca | cctctccatt | 1920 |
| tttgccttgg | tgctcatctt | aatggcctaa | tgcaccccca | aacatggaaa | tatcaccaaa | 1980 |
| aaatacttaa | tagtccacca | aaaggcaaga | ctgcccttag | aaattctagc | ctggtttgga | 2040 |
| gatactaact | gctctcagag | aaagtagctt | tgtgacatgt | catgaaccca | tgtttgcaat | 2100 |
| caaagatgat | aaaatagatt | cttattttc | ccccaccccc | gaaaatgttc | aataatgtcc | 2160 |
| catgtaaaac | ctgctacaaa | tggcagctta | tacatagcaa | tggtaaaatc | atcatctgga | 2220 |
| tttaggaatt | gctcttgtca | tacccccaag | tttctaagat | ttaagattct | ccttactact | 2280 |
| atcctacgtt | taaatatctt | tgaaagtttg | tattaaatgt | gaattttaag | aaataatatt | 2340 |
| tatatttctg | taaatgtaaa | ctgtgaagat | agttataaac | tgaagcagat | acctggaacc | 2400 |
| acctaaagaa | cttccattta | tggaggattt | ttttgcccct | tgtgtttgga | attataaaat | 2460 |
| ataggtaaaa | gtacgtaatt | aaataatgtt | tttggtaaaa | aaaaaaaaa | aaaaaaaaaa | 2520 |
| aaaaaaaaa | aaaaaaaaa | aaaaaaaaa | aaaaaaaaaa | aaa | | 2563 |
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<210> 132
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<211> 2445

<212> DNA

 $<\!\!213\!\!>$ NM_152876.1| Homo sapiens tumor necrosis factor receptor superfamily, member 6 (TNFRSF6), transcript variant 6, mRNA

| <400> 132 cctacccgcg | cgcaggccaa | gttgctgaat | caatggagcc | ctccccaacc | cgggcgttcc | 6 O |
|----------------------|------------|------------|------------|------------|------------|-------------|
| ccagcgaggc | ttccttccca | tcctcctgac | caccggggct | tttcgtgagc | tcgtctctga | 12 O |
| tctcgcgcaa | gagtgacaca | caggtgttca | aagacgcttc | tggggagtga | gggaagcggt | 18 O |
| ttacgagtga | cttggctgga | gcctcagggg | cgggcactgg | cacggaacac | accctgaggc | 24 O |
| cagccctggc | tgcccaggcg | gagctgcctc | ttctcccgcg | ggttggtgga | cccgctcagt | 30 O |
| acggagttgg | ggaagctctt | tcacttcgga | ggattgctca | acaaccatgc | tgggcatctg | 36 O |
| gaccctccta | cctctggttc | ttacgtctgt | tgctagatta | tcgtccaaaa | gtgttaatgc | 42 O |
| ccaagtgact | gacatcaact | ccaagggatt | ggaattgagg | aagactgtta | ctacagttga | 48 O |
| gactcagaac | ttggaaggcc | tgcatcatga | tggccaattc | tgccataagc | cctgtcctcc | 54 O |
| agatgtgaac | atggaatcat | caaggaatgc | acactcacca | gcaacaccaa | gtgcaaagag | 60 O |
| gaagtgaaga | gaaaggaagt | acagaaaaca | tgcagaaagc | acagaaagga | aaaccaaggt | 66 O |
| tctcatgaat | ctccaacctt | aaatcctgaa | acagtggcaa | taaatttatc | tgatgttgac | 72 0 |
| ttgagtaaat | atatcaccac | tattgctgga | gtcatgacac | taagtcaagt | taaaggcttt | 78 O |
| gttcgaaaga | atggtgtcaa | tgaagccaaa | atagatgaga | tcaagaatga | caatgtccaa | 840 |
| gacacagcag | aacagaaagt | tcaactgctt | cgtaattggc | atcaacttca | tggaaagaaa | 900 |
| gaagcgtatg | acacattgat | taaagatctc | aaaaaagcca | atctttgtac | tcttgcagag | 960 |
| aaaattcaga | ctatcatcct | caaggacatt | actagtgact | cagaaaattc | aaacttcaga | 1020 |
| aatgaaatcc | aaagcttggt | ctagagtgaa | aaacaacaaa | ttcagttctg | agtatatgca | 1080 |
| attagtgttt | gaaaagattc | ttaatagctg | gctgtaaata | ctgcttggtt | ttttactggg | 1140 |
| tacattttat | catttattag | cgctgaagag | ccaacatatt | tgtagatttt | taatatctca | 1200 |
| tgattctgcc | tccaaggatg | tttaaaatct | agttgggaaa | acaaacttca | tcaagagtaa | 1260 |
| atgcagtggc | atgctaagta | cccaaatagg | agtgtatgca | gaggatgaaa | gattaagatt | 1320 |
| atgctctggc | atctaacata | tgattctgta | gtatgaatgt | aatcagtgta | tgttagtaca | 1380 |
| aatgtctatc | cacaggctaa | ccccactcta | tgaatcaata | gaagaagcta | tgaccttttg | 1440 |
| ctgaaatatc | agttactgaa | caggcaggcc | actttgcctc | taaattacct | ctgataattc | 1500 |
| tagagatttt | accatatttc | taaactttgt | ttataactct | gagaagatca | tatttatgta | 1560 |
| aagtatatgt | atttgagtgc | agaatttaaa | taaggctcta | cctcaaagac | ctttgcacag | 1620 |
| tttattggtg | tcatattata | caatatttca | attgtgaatt | cacatagaaa | acattaaatt | 1680 |
| | | | | | | |

| ataatgtttg | actattatat | atgtgtatgc | attttactgg | ctcaaaacta | cctacttctt | 1740 |
|------------|------------|------------|------------|------------|------------|------|
| tctcaggcat | caaaagcatt | ttgagcagga | gagtattact | agagctttgc | cacctctcca | 1800 |
| tttttgcctt | ggtgctcatc | ttaatggcct | aatgcacccc | caaacatgga | aatatcacca | 1860 |
| aaaaatactt | aatagtccac | caaaaggcaa | gactgccctt | agaaattcta | gcctggtttg | 1920 |
| gagatactaa | ctgctctcag | agaaagtagc | tttgtgacat | gtcatgaacc | catgtttgca | 1980 |
| atcaaagatg | ataaaataga | ttcttatttt | tccccaccc | ccgaaaatgt | tcaataatgt | 2040 |
| cccatgtaaa | acctgctaca | aatggcagct | tatacatagc | aatggtaaaa | tcatcatctg | 2100 |
| gatttaggaa | ttgctcttgt | cataccccca | agtttctaag | atttaagatt | ctccttacta | 2160 |
| ctatcctacg | tttaaatatc | tttgaaagtt | tgtattaaat | gtgaatttta | agaaataata | 2220 |
| tttatatttc | tgtaaatgta | aactgtgaag | atagttataa | actgaagcag | atacctggaa | 2280 |
| ccacctaaag | aacttccatt | tatggaggat | ttttttgccc | cttgtgtttg | gaattataaa | 2340 |
| atataggtaa | aagtacgtaa | ttaaataatg | tttttggtaa | aaaaaaaaa | aaaaaaaaaa | 2400 |
| aaaaaaaaa | aaaaaaaaa | aaaaaaaaa | aaaaaaaaa | aaaaa | | 2445 |
| | | | | | | |

<210> 133

<211> 2508

<212> DNA

<213> NM_152877.1| Homo sapiens tumor necrosis factor receptor superfamily, member 6 (TNFRSF6), transcript variant 7, mRNA

| <400> 133 cctacccgcg | cgcaggccaa | gttgctgaat | caatggagcc | ctccccaacc | cgggcgttcc | 60 |
|----------------------|------------|------------|------------|------------|------------|-----|
| ccagcgaggc | ttccttccca | tcctcctgac | caccggggct | tttcgtgagc | tcgtctctga | 120 |
| tctcgcgcaa | gagtgacaca | caggtgttca | aagacgcttc | tggggagtga | gggaagcggt | 180 |
| ttacgagtga | cttggctgga | gcctcagggg | cgggcactgg | cacggaacac | accctgaggc | 240 |
| cagccctggc | tgcccaggcg | gagctgcctc | ttctcccgcg | ggttggtgga | cccgctcagt | 300 |
| acggagttgg | ggaagctctt | tcacttcgga | ggattgctca | acaaccatgc | tgggcatctg | 360 |
| gaccctccta | cctctggttc | ttacgtctgt | tgctagatta | tcgtccaaaa | gtgttaatgc | 420 |
| ccaagtgact | gacatcaact | ccaagggatt | ggaattgagg | aagactgtta | ctacagttga | 480 |
| gactcagaac | ttggaaggcc | tgcatcatga | tggccaattc | tgccataagc | cctgtcctcc | 540 |
| agatgtgaac | atggaatcat | caaggaatgc | acactcacca | gcaacaccaa | gtgcaaagag | 600 |
| gaaggatcca | gatctaactt | ggggtggctt | tgtcttcttc | ttttgccaat | tccactaatt | 660 |
| gtttgggtga | agagaaagga | agtacagaaa | acatgcagaa | agcacagaaa | ggaaaaccaa | 720 |
| ggttctcatg | aatctccaac | cttaaatcct | gaaacagtgg | caataaattt | atctgatgtt | 780 |
| gacttgagta | aatatatcac | cactattgct | ggagtcatga | cactaagtca | agttaaaggc | 840 |

| tttgttcgaa | agaatggtgt | caatgaagcc | aaaatagatg | agatcaagaa | tgacaatgtc | 900 |
|------------|------------|------------|------------|------------|---------------------|------|
| caagacacag | cagaacagaa | agttcaactg | cttcgtaatt | ggcatcaact | tcatg g aaag | 960 |
| aaagaagcgt | atgacacatt | gattaaagat | ctcaaaaaag | ccaatctttg | tactcttgca | 1020 |
| gagaaaattc | agactatcat | cctcaaggac | attactagtg | actcagaaaa | ttcaaacttc | 1080 |
| agaaatgaaa | tccaaagctt | ggtctagagt | gaaaaacaac | aaattcagtt | ctgagtatat | 1140 |
| gcaattagtg | tttgaaaaga | ttcttaatag | ctggctgtaa | atactgcttg | gttttttact | 1200 |
| gggtacattt | tatcatttat | tagcgctgaa | gagccaacat | atttgtagat | ttttaatatc | 1260 |
| tcatgattct | gcctccaagg | atgtttaaaa | tctagttggg | aaaacaaact | tcatcaagag | 1320 |
| taaatgcagt | ggcatgctaa | gtacccaaat | aggagtgtat | gcagaggatg | aaagattaag | 1380 |
| attatgctct | ggcatctaac | atatgattct | gtagtatgaa | tgtaatcagt | gtatgttagt | 1440 |
| acaaatgtct | atccacaggc | taaccccact | ctatgaatca | atagaagaag | ctatgacctt | 1500 |
| ttgctgaaat | atcagttact | gaacaggcag | gccactttgc | ctctaaatta | cctctgataa | 1560 |
| ttctagagat | tttaccatat | ttctaaactt | tgtttataac | tctgagaaga | tcatatttat | 1620 |
| gtaaagtata | tgtatttgag | tgcagaattt | aaataaggct | ctacctcaaa | gacctttgca | 1680 |
| cagtttattg | gtgtcatatt | atacaatatt | tcaattgtga | attcacatag | aaaacattaa | 1740 |
| attataatgt | ttgactatta | tatatgtgta | tgcattttac | tggctcaaaa | ctacctactt | 1800 |
| ctttctcagg | catcaaaagc | attttgagca | ggagagtatt | actagagctt | tgccacctct | 1860 |
| ccatttttgc | cttggtgctc | atcttaatgg | cctaatgcac | ccccaaacat | ggaaatatca | 1920 |
| ccaaaaaata | cttaatagtc | caccaaaagg | caagactgcc | cttagaaatt | ctagcctggt | 1980 |
| ttggagatac | taactgctct | cagagaaagt | agctttgtga | catgtcatga | acccatgttt | 2040 |
| gcaatcaaag | atgataaaat | agattcttat | ttttccccca | ccccgaaaa | tgttcaataa | 2100 |
| tgtcccatgt | aaaacctgct | acaaatggca | gcttatacat | agcaatggta | aaatcatcat | 2160 |
| ctggatttag | gaattgctct | tgtcataccc | ccaagtttct | aagatttaag | attctcctta | 2220 |
| ctactatcct | acgtttaaat | atctttgaaa | gtttgtatta | aatgtgaatt | ttaagaaata | 2280 |
| atatttatat | ttctgtaaat | gtaaactgtg | aagatagtta | taaactgaag | cagatacctg | 2340 |
| gaaccaccta | aagaacttcc | atttatggag | gattttttg | ccccttgtgt | ttggaattat | 2400 |
| aaaatatagg | taaaagtacg | taattaaata | atgtttttgg | taaaaaaaaa | aaaaaaaaa | 2460 |
| aaaaaaaaa | aaaaaaaaa | aaaaaaaaa | aaaaaaaaa | aaaaaaaa | | 2508 |
| | | | | | | |

<210> 134

<211> 2583

<212> DNA

 $<\!\!213\!\!>$ NM_152875.1| Homo sapiens tumor necrosis factor receptor superfamily, member 6 (TNFRSF6), transcript variant 5, mRNA

| <400> 134 cctacccgcg | cgcaggccaa | gttgctgaat | caatggagcc | ctccccaacc | cgggcgttcc | 60 |
|----------------------|------------|------------|------------|---------------------|------------|------|
| ccagcgaggc | ttccttccca | tcctcctgac | caccggggct | tttcgtgagc | tcgtctctga | 120 |
| tctcgcgcaa | gagtgacaca | caggtgttca | aagacgcttc | tggggagtga | gggaagcggt | 180 |
| ttacgagtga | cttggctgga | gcctcagggg | cgggcactgg | cacggaacac | accctgaggc | 240 |
| cagccctggc | tgcccaggcg | gagctgcctc | ttctcccgcg | ggttggtgga | cccgctcagt | 300 |
| acggagttgg | ggaagctctt | tcacttcgga | ggattgctca | acaac catgc | tgggcatctg | 360 |
| gaccctccta | cctctggttc | ttacgtctgt | tgctagatta | tcgtccaaaa | gtgttaatgc | 420 |
| ccaagtgact | gacatcaact | ccaagggatt | ggaattgagg | aagactgtta | ctacagttga | 480 |
| gactcagaac | ttggaaggcc | tgcatcatga | tggccaattc | tgccataagc | cctgtcctcc | 540 |
| aggtgaaagg | aaagctaggg | actgcacagt | caatggggat | gaaccagact | gcgtgccctg | 600 |
| ccaagaaggg | aaggagtaca | cagacaaagc | ccatttttct | tccaaatgca | gaagatgtag | 660 |
| attgtgtgat | gaaggacatg | atgtgaacat | ggaatcatca | aggaatgcac | actcaccagc | 720 |
| aacaccaagt | gcaaagagga | agtgaagaga | aaggaagtac | agaaaacatg | cagaaagcac | 780 |
| agaaaggaaa | accaaggttc | tcatgaatct | ccaaccttaa | atcctgaaac | agtggcaata | 840 |
| aatttatctg | atgttgactt | gagtaaatat | atcaccacta | ttgctggagt | catgacacta | 900 |
| agtcaagtta | aaggctttgt | tcgaaagaat | ggtgtcaatg | aagccaaaat | agatgagatc | 960 |
| aagaatgaca | atgtccaaga | cacagcagaa | cagaaagttc | aactgcttcg | taattggcat | 1020 |
| caacttcatg | gaaagaaaga | agcgtatgac | acattgatta | aagat c tcaa | aaaagccaat | 1080 |
| ctttgtactc | ttgcagagaa | aattcagact | atcatcctca | aggacattac | tagtgactca | 1140 |
| gaaaattcaa | acttcagaaa | tgaaatccaa | agcttggtct | agagtgaaaa | acaacaaatt | 1200 |
| cagttctgag | tatatgcaat | tagtgtttga | aaagattctt | aatagctggc | tgtaaatact | 1260 |
| gcttggtttt | ttactgggta | cattttatca | tttattagcg | ctgaagagcc | aacatatttg | 1320 |
| tagattttta | atatctcatg | attctgcctc | caaggatgtt | taaaatctag | ttgggaaaac | 1380 |
| aaacttcatc | aagagtaaat | gcagtggcat | gctaagtacc | caaataggag | tgtatgcaga | 1440 |
| ggatgaaaga | ttaagattat | gctctggcat | ctaacatatg | attctgtagt | atgaatgtaa | 1500 |
| tcagtgtatg | ttagtacaaa | tgtctatcca | caggctaacc | ccact ctatg | aatcaataga | 1560 |
| agaagctatg | accttttgct | gaaatatcag | ttactgaaca | ggcaggccac | tttgcctcta | 1620 |
| aattacctct | gataattcta | gagattttac | catatttcta | aactttgttt | ataactctga | 1680 |
| gaagatcata | tttatgtaaa | gtatatgtat | ttgagtgcag | aatttaaata | aggctctacc | 1740 |
| tcaaagacct | ttgcacagtt | tattggtgtc | atattataca | atatttcaat | tgtgaattca | 1800 |
| catagaaaac | attaaattat | aatgtttgac | tattatatat | gtgtatgcat | tttactggct | 1860 |
| caaaactacc | tacttctttc | tcaggcatca | aaagcatttt | gagcaggaga | gtattactag | 1920 |
| | | | | | | |

| agctttgcca cctctccatt tttgccttgg tgctcatctt aatggcctaa tgcacccca | 1980 |
|---|-------------------------|
| aacatggaaa tatcaccaaa aaatacttaa tagtccacca aaaggcaaga ctgcccttag | 2040 |
| aaattctagc ctggtttgga gatactaact gctctcagag aaagtagctt tgtgacatgt | 2100 |
| catgaaccca tgtttgcaat caaagatgat aaaatagatt cttatttttc ccccacccc | 2160 |
| gaaaatgttc aataatgtcc catgtaaaac ctgctacaaa tggcagctta tacatagcaa | 2220 |
| tggtaaaatc atcatctgga tttaggaatt gctcttgtca tacccccaag tttctaagat | 2280 |
| ttaagattct ccttactact atcctacgtt taaatatctt tgaaagtttg tattaaatgt | 2340 |
| gaattttaag aaataatatt tatatttctg taaatgtaaa ctgtgaagat agttataaac | 2400 |
| tgaagcagat acctggaacc acctaaagaa cttccattta tggaggattt ttttgcccct | 2460 |
| tgtgtttgga attataaaat ataggtaaaa gtacgtaatt aaataatgtt tttggtaaaa | 2520 |
| aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaa | 2580 |
| aaa | 2583 |
| <210> 135 | |
| <211> 316 | |
| <212> DNA | |
| <213> >gi 13310411 gb AF333388.1 AF333388 Homo sapiens metallothic | onein 1H-l [.] |
| protein mRNA, complete cds | |
| | |
| <400> 135 cctcttctct tctcgcttgg gaacgccggt ctcacctcgg cttgcaatgg accccaactg | 60 |
| ctcctgcgcc gctggaggct cctacgcctg cgccggctcc tgcaagtgca aaaagtgcaa | 120 |
| atgcacctcc tgcaagaaga gctgctgctc ctgttgcccc ctgggctgtg ccaagtgtgc | 180 |
| ccagggctgc atccgcaaag gggcttcgga aaagtgcagc tgctgtgcct gatgtcggga | 240 |
| ctgccctgct ctcggatgaa aacagaatga cacgtaaagt ccgggatttt tttttctaca | 300 |
| actccgactc atttgc | 316 |
| | |
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| <211> 3145 | |
| <212> DNA | |
| <213> NM_000251. Homo sapiens muts[gi:4557760] | |
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| <400> 136 ggcgggaaac agcttagtgg gtgtggggtc gcgcattttc ttcaaccagg aggtgaggag | 60 |
| | |

ike

120

180

gtttcgacat ggcggtgcag ccgaaggaga cgctgcagtt ggagagcgcg gccgaggtcg

gcttcgtgcg cttctttcag ggcatgccgg agaagccgac caccacagtg cgccttttcg

| accggggcga | cttctatacg | gcgcacggcg | aggacgcgct | gctggccgcc | cgggaggtgt | 240 |
|------------|------------|------------|--------------------|------------|------------|------|
| tcaagaccca | gggggtgatc | aagtacatgg | ggccggcagg | agcaaagaat | ctgcagagtg | 300 |
| ttgtgcttag | taaaatgaat | tttgaatctt | ttgtaaaaga | tcttcttctg | gttcgtcagt | 360 |
| atagagttga | agtttataag | aatagagctg | gaaataagg c | atccaaggag | aatgattggt | 420 |
| atttggcata | taaggcttct | cctggcaatc | tctctcagtt | tgaagacatt | ctctttggta | 480 |
| acaatgatat | gtcagcttcc | attggtgttg | tgggtgttaa | aatgtccgca | gttgatggcc | 540 |
| agagacaggt | tggagttggg | tatgtggatt | ccatacagag | gaaactagga | ctgtgtgaat | 600 |
| tccctgataa | tgatcagttc | tccaatcttg | aggctctcct | catccagatt | ggaccaaagg | 660 |
| aatgtgtttt | acccggagga | gagactgctg | gagacatgg g | gaaactgaga | cagataattc | 720 |
| aaagaggagg | aattctgatc | acagaaagaa | aaaaagctga | cttttccaca | aaagacattt | 780 |
| atcaggacct | caaccggttg | ttgaaaggca | aaaagggaga | gcagatgaat | agtgctgtat | 840 |
| tgccagaaat | ggagaatcag | gttgcagttt | catcactgt c | tgcggtaatc | aagtttttag | 900 |
| aactcttatc | agatgattcc | aactttggac | agtttgaact | gactactttt | gacttcagcc | 960 |
| agtatatgaa | attggatatt | gcagcagtca | gagcccttaa | cctttttcag | ggttctgttg | 1020 |
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660

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